10-0.53 663

Protein Sequence Searches - February 2005

All of the sequence databases on ABSS have recomb been and red

- Please note that the curators of the UniProt database have purged some temporary accession numbers from the most recent version of UniProt. These sequences have been assigned new permanent accession numbers. The new UniProt record may not contain the previous temporary accession number.
- If you encounter an accession number from an older search run against UniProt (results file extension .rup) that can no longer be found in the database, the permanent record with the new accession number can be found by searching the old accession number in the UniProt Protein Archive database (UniPARC) at:

http://www.pir.uniprot.org/database/archive.shtml

If you have any questions regarding this information or your results, please contact any STIC searcher.

When submitting sequence search results for scanning into IFW, please include a copy of this attachment to assist any future Examiners or members of the public who may encounter UniProt temporary accession numbers.

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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on:

March 23, 2005, 14:38:05; Search time 23.5 Seconds (without alignments) 20.472 Million cell updates/sec

SEQ1 27 1 ffglm 5

Title: Perfect score: Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

283416 seqs, 96216763 residues Searched:

Total number of hits satisfying chosen parameters:

206

Minimum DB seq length: 0 Maximum DB seq length: 5

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

PIR 79:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	Ig heavy chain CRD	alkanal monooxygen	neuropeptide Antho	Met-enkephalin - b	Ig heavy chain CRD	T-cell receptor be	27.5 kda structura	27.5K structural p	27.5K structural p	28.5K structural p	28K structural pro	blood cell protein	cholecystokinin-5	photosystem I 10.4	Leu-enkephalin - b	T-cell receptor be		spinal cord_peptid	synaptosomal-assoc	T-cell receptor be	hypothetical prote	gut pentapeptide -	surface protein te	н	T-cell receptor be	growth-modulating	- chi	spinal cord peptid	RFamic
SUMMARIES	ID	PT0278	A44955	JQ1273	A61445	PT0240	A53284	G44817	144817	E44817	C44817	A44817	S68328	A32516	PQ0689	B61445	PT0633	PT0572	B23751	E44823	B53284	T10954	JH0253	S69237	PT0636	PT0571	GKHU	A60898	A23751	ECXAA
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hypothetical prote starvation-induced hypothetical prote	COI intron 16 prot autho-RF amide neu RPCH-related neuro	myosin-light-chain Ig mu chain V regi ubiquitin - rat	globulin IV alpha carbon-monoxide de neuropeptide Antho	FMRFamide - polych achatin-I - giant Ig heavy chain CRD	T-cell receptor be
D41654 S53508 T30569	138888 A25844 A34626	S39390 S43959 S47552	S09478 PL0140 A35779	A60418 A32480 PT0271	PT0711
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30 31	33.3 35.4 35.4	36 37 38	39 4 4 1	4 4 4 Çi E 4	45

ALIGNMENTS

RESULT 1 Program Ig heavy chain CRD3 region (clone 4-88) - human (fragment) C;Species: Homo sapiens (man) C;Date: 30-5ep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996 C;Accession: PP0278 R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G. J. Exp. Med. 173, 395-407, 1991 A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and jc A;Reference number: PT0228 A;Reference number: PT0228 A;Reference number: PT0228 A;Residues: 1-5 «YAM» A;Residues: 1-5 «YAM» A;Residues: 1-5 «YAM» A;Experimental source: B lymphocyte C;Reywords: heterotetramer; immunoglobulin	Query Match Best Local Similarity 40.0%; Pred. No. 2.8e+05; Matches 2; Conservative 3; Mismatches 0; Indels 0; Gaps 0; 1 FFGLM S 1
RESULT 1 PT0278 Ig heavy chu C; Species: 1 C; Accesion C; Accesion R; Yamada, M J. Exp. Med A; Title: Prr A; Reference A; Resiones: A; Resiques: A; Experiment C; Keywords:	Query Matc Best Loca: Matches Qy Db

alkanal monooxygenase (FWN-linked) (EC 1.14.14.3) alpha chain - Vibrio harveyi (fragment) C;Species: Vibrio harveyi
C;Species: Vibrio harveyi
C;Date: 03-0un-1993 #sequence_revision 03-0un-1993 #text_change 26-May-2000
C;Accession: A44955
R;Paquatte, O.; Tu, S.C.
Photochem. Photobiol. 50, 817-825, 1989
A;Title: Chemical modification and characterization of the alpha cysteine 106 at the Vibn A;Reference number: A44955; MUID:90175700; PMID:2626493
A;Reference number: A44955
A;Reture: preliminary
A;Molecule type: protein
A;Residuse: 1-5 < PRQ>
C;Keywords: FMN; luminescence; monooxygenase; oxidoreductase

Gaps ö Query Match
Best Local Similarity 66.7%; Pred. No. 2.8e+05;
Matches 2; Conservative 1; Mismatches 0; Indels

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2 FGL 4

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RESULT 3 JQ1273

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seql.rpr

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T-cell receptor beta 2 chain D region, Dbeta2 - rabbit
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Decies: O2-May-1994 #sequence_revision 18-Nov-1994 #text_change O5-Nov-1999
C;Accession: A5:284
R;Harindranath, N; Alexander, C.B.; Mage, R.G.
Mol. Immunol. 28, 881-888, 1991
A;ritle: Evolutionarily conserved organization and sequences of germline diversity and journalist or minor. A;Reference number a5:3284; MuID:91342695; PMID:1678859
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C;Species: Leuconostoc oenos phage P32
C;Bate: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
C;Accession: G44811,
J; Lonvaud, A.; Hammes, W.P.
J; Gen. Microbiol. 137, 2135-2139, 1991
A;Title: Lysogeny in Leuconostoc oenos.
A;Reference number: A44817; MUID:92085033; PMID:1748868
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Cibate: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
CiAccession: 144817
CiAccession: 144817
Airlie: Lysogeny in Leuconostoc oenos.
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Best Local Similarity 100.0%; Score 10; DB 2; Length 5;
Matches 2; Conservative 0; Mismatches 0; Indels
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A.Residues: 1-5 <ARE>
A.Note: sequence extracted from NCBI backbone (NCBIP:70333)
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Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 2; Conservative 0; Mismatches 0;
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                                             C. Accession: JO273

R. Mar-1992 #sequence_revision 04-Dec-1992 #text_change 09-Jul-2004

C. Accession: JO273

R. Mochen. Biophys. Res. Commun. 179, 1205-1211, 1991

R. Mille: Isolation of L.-3-phenyllactyl-phe-1993 Multi-1681803

A. Mille: Isolation of L.-3-phenyllactyl-phe-1995 Ala-NH2 (Antho-Kaamide), a novel neuropep A. Accession: JO1273

R. Molecule type: protein

R. Molecule type: protein

R. Comment: The carboxyl-terminal amide probably arises from cleavage of a following glyctyl Kassidues: 1-4 kNOT2

C. Comment: The carboxyl-terminal amide probably arises from cleavage of a following glyctyl Kassidues: 1-4 kNOT3

R. Modified site: L-3-phenyllactic acid (Phe) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Met-enkephalin - blue mussel
C;Species: Mytilus edulis (blue mussel)
C;Scession: A61445
R;Leung, M.K.; Stefano, G.B.
R;Revence number: A61445; MUID:84144823; PMID:6583690
A;Recession: A61445
R;Molecule type: protein
A;Residues: 1-5 c.EUJ
A;Residues: 1-5 c.EUJ
A;Residues: 1-5 c.EUJ
A;Residues: neuropeptide; opioid peptide
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19 heavy chain CRD3 region (clone 2-100B) - human (fragment)

C; Date: 30-Sep-1933 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996

C; Accession: PT0240

R; Yamada, M; Wasserman, R; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.

Exp. Med. 173, 395-407, 1991

A; Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A; Reference number: PT0222; MUID:91108337; PMID:1899102
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                          Species: Anthopleura elegantissima
Date: 31-Mar-1992 #sequence_revision 04-Dec-1992 #text_change 09-Jul-2004
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neuropeptide Antho-KAamide - sea anemone (Anthopleura elegantissima)
C;Species: Anthopleura elegantissima
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ches 0; Indels
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37.0%; Score 10; DB 2; Length 4;
Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 2; Conservative 0; Mismatches 0; Indels
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44.4%; Score 12; DB :
Best Local Similarity 100.0%; Pred. No. 2.8
Matches 2; Conservative 0; Mismatches
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A;Residues: 1-4 <YAM>
A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; lmmunoglobulin
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Best Local Similarity 66.7
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Cholecystokinin-5 - dog
NyAlternate names: CCK-5
Cippedies: Canis lupus familiaris (dog)
Cjaccession: A32516
RyShively, J.; Reeve Jr., J.R.; Eysselein, V.E.; Ben-Avram, C.; Vigna, S.R.; Walsh, J.H.
An. J. Physiol. 252, G272-G275, 1987
An. J. Physiol. 252, G272-G275, 1987
An. J. Physiol. 252, G272-G275, 1987
An. Streeve Jr., J.R.; Eysselein, V.E.; Ben-Avram, C.; Vigna, S.R.; Walsh, J.H.
A, Fittle: CCK-5: sequence analysis of a small cholecystokinin from canine brain and intest
A, Fittle: CCK-5: sequence analysis of a small cholecystokinin from canine brain and intest
A, Fittle: CCK-5: sequence analysis of the five carboxyl-terminal residues of cholecystok
C, Comment: This peptide corresponds to the five carboxyl-terminal residues of cholecystok
C, Superfamily: gastrin
C, Goment: This peptide carboxyl end; neuropeptide
C, Superfamily: gastrin
C, Goment: This peptide carboxyl end (Phe) #status experimental
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C;Species: Nicotiana tabacum (common tobacco)
C;Species: Nicotiana tabacum (common tobacco)
C;Species: Nicotiana tabacum (common tobacco)
C;Species: 19-Way-1994 #sequence_revision 19-Way-1994 #text_change 17-Mar-1999
C;Acces: 10-Way-1994 #sequence_revision 19-Way-1994 #text_change 17-Mar-1999
R;Obokata, J.; Mikami, K.; Hayashida, N.; Nakamura, M.; Sugiura, M.
Plant Physiol. 102, 1259-1267, 1993
A;Title: Molecular heterogeneity of photosystem I. psaD, psaE, psaF, psaH and psaL are all A;Reference number: PQ0667; MUID:94105345; PMID:8278548
A;Accession: PQ0689
A;Molecule type: protein
A;Residues: 1-5 <0BO>
C;Keywords: chloroplast; photosynthesis; photosystem I; thylakoid
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50.0%; Pred. No. 2.8e+05;
iive 1; Mismatches 0
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RESULT 9
E44817
27.5K structural protein - Leuconostoc oenos phage P54 (fragment)
C.Species: Leuconostoc oenos phage P54
C.Bete: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
C.Accession: B44817
C.Accessi
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C;Species: Leuconostoc oenos phage PAt5-12
C;Accession: C44817
R;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J, Gen. Microbiol. 137, 2135-2139, 1991
A;Title: Lysogeny in Leuconostoc oenos.
A;Reference number: A44817; MUID:92085033; PMID:1748868
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C;Species: Leuconostoc oenos phage PZtll-15
C;Adzes: J-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
C;Adzession: A44817
R;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J. Gen. Microbiol. 137, 2135-2139, 1991
A;Title: Lysogeny in Leuconostoc oenos.
A;Reference number: A44817; MUID:92085033; PMID:1748868
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A;Residues: 1-5 <ARE>
A;Note: sequence extracted from NCBI backbone (NCBIP:70341)
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A;Residues: 1-5 <ARE>
A;Note: sequence extracted from NCBI backbone (NCBIP:70343)
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Wed Mar 23 15:33:42 2005
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Leu-enkephalin - blue mussel
C;Species: Mytilus edulis (blue mussel)
C;Accession: B61445
Proc. Natl. Acad. Sci. U.S.A. 81, 955-958, 1984
A;Title: Isolation and identification of enkephalins in pedal ganglia of Mytilus edulis
A;Reference number: A61445; MUD:84144823; PMID:6583690
A;Molecule type: protein
A;Residues: 1-5 clEUs
A;Residues: 1-5 clEUs
C;Keywords: neuropeptide; opioid peptide
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Search completed: March 23, 2005, 14:51:53 Job time : 25.5 secs

GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

March 23, 2005, 14:21:29 ; Search time 112.5 Seconds (without alignments) 22.759 Million cell updates/sec Run on:

27 1 ffglm 5 Perfect score: Sequence:

BLOSUM62 Scoring table:

Gapop 10.0 , Gapext 0.5

1612378 segs, 512079187 residues Searched: Total number of hits satisfying chosen parameters:

23

Minimum DB seq length: 0 Maximum DB seq length: 5

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

UniProt_03:*
1: uniprot_sprot:*
2: uniprot_trembl:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result	_	* Query			SOFIFERED		
Score	_ ;	atch	Match Length	BB	QI	Description	a .
		59.3	S	-	AL14 CARMA		carcinus ma
-		44.4	4	Н	FFKA ANTEL	'n	anthopleura
٦		44.4	4	Н	OCP1_OCTMI		octopus min
-	2	44.4	Ŋ	Н	PAP2_PARMA	P81864 parc	pardachirus
-		44.4	2	Н	RE11_LITRU	P82070 lite	litoria rub
-	-	44.4	5	Н	RE21_LITRU	P82071 lite	litoria rub
1		44.4	2	Н	RE31_LITRU	P82072 lite	litoria rub
1	2	14.4	2	Н	RE32_LITRU	P82073 litc	litoria rub
1	2	44.4	S	Н	UC22_MAIZE	P80628 zea	mays (m
7	1 4	40.7	ū	-	TPIS_CANFA	P54714 cani	canis famil
	6	33.3	4	Н	FYRI ANTEL		anthopleura
	6	33.3	4	-	ILME_SEPOF		ia offic
		25.9	2	Н	UF01 MOUSE	σ	musculu
	9	22.2	7	Н	GWA SEPOF	P83570 sepia	ia offic
	.,	22.2	c	Н	GRWM HUMAN	P01157 homo	o sapien
	9	22.2	4	Н	ACH1_ACHFU	-	achatina fu
	.,		4	Н	DCML PSECH		pseudomonas
	.,	22.2	4	Н	EOSI HUMAN	P02731 homo	homo sapien
			4	Н	FAR3_HIRME	~	nirudo medi
		22.2	4	-	FAR4 HIRME	_	hirudo medi
		22.2	4	Н	FLRF_HIRME		hirudo medi
		22.2	4	-	FLRN ANTEL	P58707 ant	anthopleura
		22.2	4	Н	FMRF MACNI		macrocallis
			4	~	OCP3_OCTMI	-	octopus min
		22.2	4	~	Q160 <u>4</u> 7	Q16047 homo	o sapien
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			ß	7	E103_LITRU	P82099 lite	litoria rub
			ιΩ	7	EI04_LITRU		litoria rub
	9		S	Н	FARP ARTTR	~	artioposthi
		22.2	S	-	FARP_CHICK	3308	lus gall
	9	22.2	ស	Н	SUGA_ACHDO	P19991 acheta	eta dome

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Gaps

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59.3%; Score 16; DB 1; Length 5; 100.0%; Pred. No. 1.6e+06; Live 0; Mismatches 0; Indels

3; Conservative

FGL 5 2 FGL 4

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Query Match Best Local Similarity Matches 3; Conserv

MEDLINE=92028852; PubMed=1681803; Nothacker H.-P., Rinehart K.L. Jr., Grimmelikhuijzen C.J.P.; "Isolation of L-3-phenyllactyl-Phe-Lys-Ala-NH2 (Antho-KAamide), a

SEQUENCE

PERSULT 2

PERA ANTEL

D PERA ANTEL

STANDARD; PRT; 4 AA.

PST05;

DT 28-FEB-2003 (Rel. 41, Created)

DT 28-FEB-2003 (Rel. 44, Last sequence update)

DT 05-UTL-2004 (Rel. 44, Last annotation update)

B Antho-KAamide.

OS Anthopleura elegantissima (Sea anemone).

OC Eukaryota; Metazoa; Cnidaria; Anthozoa; Zoantharia; Actiniaria;

OX Nymantheae; Actiniidae; Anthopleura.

P38005 chlamydia t P19918 pseudomonas Q96at0 homo sapien P13071 citrobacter P12997 citrobacter Q99007 hordeum vul P83073 bacillus ce Q08433 rattus sp. P67859 limulus pol P67859 limulus pol P67859 periplaneta P58261 daucus caro P24272 vibrio fisc P36515 saccharomyc	T.1 CARMA T.14 CARMA STANDARD; PRT; 5 AA. PR114 CARMA STANDARD; PRT; 5 AA. PR115 CARMA STANDARD; PRT; 5 AA. PR1817; So-MAY-2000 (Rel. 39, Last sequence update) So-MAY-2000 (Rel. 39, Last sequence update) So-MAY-2000 (Rel. 39, Last sequence update) So-Carcinus maenas (Common shore crab) (Green crab). Carcinus maenas (Common shore crab) (Green crab). Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca; Eukaryota; Metazoa; Arthropoda; Pleocyemata; Brachyura; Eukaryota; Metazoa; Arthropoda; Pleocyemata; Brachyura; Eukaryota; Metazoa; Arthropoda; Portunidae; Carcinus. NOBI_TAXID=6759; [1] TISSUE-cerebral ganglion, and Thoracic ganglion; MEDLINE-98121193; PubMed=9461295; MEDLINE-98121193; PubMed=9461295; TISSUE-cerebral ganglion, and the shore crab Carcinus maenas."; Thorpe A.; Johnsen A.H., Maestro JL., Scott A.G., Jaros P.P., Thorpe A.; Johnsen 250:727-734(1997)
UXA4 CHLTR DCMS_PSECH Q96AT0 BIOA_CITFR BIOB_CITFR 009907 Q08433 RCT_CARMA PRCT_LIMPO PR	ALIGNMENTS PRT; 5 AA. tt sequence update) tt annotation update) thore crab) (Green crab) Opda; Crustacea; Malac. Decapoda; Pleocyemata; Portunidae; Carcinus. Portunidae; Carcinus. and Thoracic ganglion; 9461295; testro JL., Scott A.G. ttion of multiple neurop in the shore crab Carcin in the shore crab Carcin on the allatostatin fami sequencing; Multipne Leucine amide (Pote
22.2 18.5 18.5 18.5 18.5 18.5 18.5 14.8 14.8 14.8 11.1 11.1 3.7 4 4 4 1 11.1 11.1 3.7	T.1 CARMA ALIACARMA STANDARD; PRT; 5 AA. ALIACARMA STANDARD; PRT; 5 AA. 30-MAY-2000 (Rel. 39, Created) 30-MAY-2000 (Rel. 39, Last sequence update) Carcinustatin 14. Carcinus maenas (Common shore crab) (Green crab) Carcinus maenas (Common shore crab) (Green crab) Eukaryota; Metazoa; Arthropoda; Crustacea; Malac Eukaracostraca; Eucarida; Decapoda; Pleocyemata; Eubrachyura; Portunoidea; Portunidae; Carcinus. NCBL TaxID=6759; [1] SEQUENCE TISSUENCE TO THOR A. Maestro JL., Scott A.G. Thorpe A.; Namestro JL., Scott A.G. Thorpe A.; Namestro JL., Scott A.G. Thorpe A.; Daiochain superfamily in the shore crab Carcinal allacostatin superfamily in the shore crab Carcinal Latostatin superfamily in the shore crab Carcinal Latostatin Sequencing; Multigene MOD LES S. S. Leucine amide (Poce SEQUENCE S. AA; S86 MW; 67287995AB300000 CRC6
3332 33333 3444 5443 5443 55443 57444 57444 675 675 675 675 675 675 675 675 675 675	ARWA 114 CARWA 1154 CARWA 1154 CARWA 1154 CARWA 1156 CO MAY-2000 10 CMAY-2000 10 CMAY-2000 10 CMAY-2000 10 CMAY-2000 11 CMAY-2000 11 CMAY-2000 12 CMAY-2000 12 CMAY-2000 13 CMAY-2000 14 CMAY-2000 15 CMAY-2000 15 CMAY-2000 15 CMAY-2000 15 CMAY-2000 15 CMAY-2000 16 CMAY-2000 17 CM
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SEQUENCE.
TISSUE-Skin secretion,
MEDLINE-Skin secretion,
Lazarovici P., Primox N., Loew L.M.;
"Purification and pore-forming activity of two hydrophobic polyreptides from the secretion of the Red sea moses sole (Pardachirus
                                                                                                  Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi, Actinopterygii, Neopterygii, Teleostei, Buteleostei, Neoteleostei, Soleoidei, Soleidae, Pardachirus.

NCBI_TaxID=31087;
                                                                                                                                                                                                                                                                                                                          ### And Properties. At high concentration extends on the sea mouse buse transacon J. Biol. Chem. 261:16704-16713(1986).

-!- FUNCTION: Exhibite unusual shark repellent and surfactant properties. Porms voltage-dependent, ion-permeable channels in membranes. At high concentration causes cell membrane lysis.
-!- SUBCELIGIAR LOCATION: Secreted.
-!- SINDIARITY: Belongs to the pardaxin family.
Direct protein sequencing; Toxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
26-JUL-2004 (Rel. 44, Last annotation update)
Rubellidin 1.1.
Litoria rubella (Desert tree frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea; Hylidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE, AND MASS SPECTROMETRY.
STSGUE-Skin secretion;
Steinborner S.T., Wabnitz P.A., Waugh R.J., Bowie J.H., Gao C.,
Tyler M.J., Wallace J.C.;
Tyler M.J., Wallace J.C.;
Tyler M.J., Wallace J.C.;
The structure of new peptides from the Australin red tree frog of evolutionary trends of amphibians.";
Litoria rubella. The skin peptide profile as a probe for the st Aust. J. Chem. 49:955-963 (1996).
-!- FINCT Chem. 49:955-963 (1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  activity.
--- SUBCELLULAR LOCATION: Secreted.
--- TISSUB SPECIFCITY: Expressed by the skin dorsal glands.
--- MASS SPECTROMETRY: MW=598; METHOD=FAB; RANGE=1-5; NOTE=Ref.1.
Amphibian defense pepilde; Direct protein sequencing.
SEQUENCE 5 AA; 598 MW; 6DD9C9CAB2A00000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        44.4%; Score 12; DB 1; Length 5; 100.0%; Pred. No. 1.6e+06; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5 AA; 614 MW; 7769C9C9C8100000 CRC64;
                               30-MAY-2000 (Rel. 39, Last sequence update)
05-UUL-2004 (Rel. 44, Last annotation update)
Pardaxin II (PXII) (Fragment)
Pardachirus marmoratus (Red sea moses sole)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 100.
Matches 2; Conservative
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Best Local Similarity 100.
Matches 2; Conservative
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ID REIL LITRU
AC P82070;
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                                                                   MEDLINE-93391436; PubMed=8397415;
McFarlane I.D., Hudman D., Nothacker H.-P., Grimmelikhuijzen C.J.P.;
The expansion behaviour of sea anemones may be coordinated by two inhibitory neuropeptides, Antho-KAamide and Antho-RIamide by two Proc. R. Soc. Lond. B. Biol. Sci. 253:183-1881993).
-I- FUNCTION: Inhibits spontaneous contractions in several muscle groups. May be involved in the expansion phase of feeding behaviour in sea anemones.
                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUE=Brain,
MEDLINE=20336815, PubMed=10876044, DOI=10.1016/S0196-9781(00)00201-1;
Wakoshi E., Hisada M., Minakara H.,
"Cardioactive peptides isolated from the brain of a Japanese octopus,
Octopus minor.",
Peptides 21:623-630(2000).
Peptides 21:623-630(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -!- SUBCELLULAR LOCATION: Secreted.
-!- PIM: Ocp-2 has L-Phe instead of D-Phe.
-!- MASS SPECTROMETRY: MW=395.2; METHOD=MALDI; RANGE=1-4; NOTE=Ref.1.
MOD RES
SEQUENCE 4 AA; 394 MW; 6AA879C810000000 CRC64;
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98-FEB-2003 (Rel. 41, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
cardioactive peptides Ocp-1/Ocp-2.
Cutogus minor (Octopus).
Eukaryota; Metazoa; Mollusca; Cephalopoda; Coleoidea; Neocoleoidea;
Octopodiformes; Octopoda; Incirrata; Octopodidae; Octopus.
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                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
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                                                                                                                                                                                                 ovel neuropeptide from sea anemones.";
iochem. Biophys. Res. Commun. 179:1205-1211(1991)
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P81864;
30-MAY-2000 (Rel. 39, Created)
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Matches 2; Conservative
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TISSUE=Skin secretion;
Query Match
Best Local Similarity
These 2; Conserva
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P82073;
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P80628;
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Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea; Hylidae;
Pelodryadinae; Litoria.
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-!- TISSUB SPECIFICITY: Expressed by the skin dorsal glands.
-!- MASS SPECTROMETRY: MW=655, METHOD=FAB; RANGE=1-5; NOTE=Ref.1.
Amidation; Amphibian defense peptide; Direct protein sequencing.
MOD RES 5 Threonine amide.
SEQÜENCE 5 AA; 656 MW; 71A9C9CB10300000 CRC64;
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-!- TISSUE SPECIFICITY: Expressed by the skin dorsal glands.
-!- TAISSUE SPECIFICITY: Expressed by the skin dorsal glands.
-!- MASS SPECIFORMETRY: MW=626, MR=HOD=FRAB, RANGE=1-5, NOTE=Ref.1.
Amphibian defense peptide, Direct protein sequencing.
SEQUENCE 5 AA; 626 MW; 6DD9C9CB10300000 CRC64;
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Steinborner S.T., Wabnitz P.A., Waugh R.J., Bowie J.H., Gao C.,
Tyler M.J., Wallace J.C.;
"The structure of new peptides from the Australin red tree frog
'Intoria rubella'. The skin peptide profile as a probe for the st
of evolutionary trends of amphibians.";
Aust. J. Chem. 49:1955-963 (1996).
-I- FUNCTION: Shows neither neuropeptide activity nor antibiotic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Aust. J. Chem. 49:955-963(1996).
-!- FUNCTION: Shows neither neuropeptide activity nor antibiotic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Steinborner S.T., Wabnitz P.A., Waugh R.J., Bowie J.H., Gao C., Tyler M.J., Wallace J.C.; "The structure of new peptides from the Australin red tree frog 'Litoria rubella'. The skin peptide profile as a probe for the sof
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                                                                                                                                                                                                                                                                                                       28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Rubellidin 2.1.
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(Rel. 41, Last sequence update)
(Rel. 44, Last annotation update)
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Litoria rubella (Desert tree frog)
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DT 08-JUL-2004

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P82072;
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RE31 LITRU

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Pernollet J.-C., Zivy M., de Vienne D.;
"The maize two dimensional gel protein database: towards an integrated
genome analysis program.";
Theor. Appl. Genet. 93:997-1005(1996).
-!- MISCELLANBOUS: On the 2D-gel the determined pI of this unknown
protein is: 6.1, its MW is: 30.4 kba.
Maize-2DPAGE; PRO628; COLEOPTILE.
MaizebB; 123954; -.
Direct protein sequencing.
                                                                Gape
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Litori electrica. Comparison with the skin peptides from Litoria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae,
PACCAD clade, Panicoideae, Andropogoneae, Zea.
                                                                                                                                                                                                                                                                                                                                                                                              28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Rubellidin 3.2.
Litoria rubella (Desert tree frog).
Litoria rubella (Desert Saniata, Vertebrata, Euteleostomi, Amphibia; Batrachia, Anura; Neobatrachia; Hyloidea; Hylidae;
Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rubella.";
Aust. J. Chem. 52:639-645(1999).
-!- FUNCTION: Shows neither neuropeptide activity nor antibiotic
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01-0cT-1996 (Rel. 34, Last sequence update)
05-UUL-2004 (Rel. 44, Last annotation update)
Uhknown protein from 2D-PAGE of etiolated coleoptile (Spot 474)
(Fragment)
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-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Expressed by the skin dorsal glands.
-!- TISSUE SPECIFICITY: Expressed by the skin dorsal glands.
-!- Amphibian defense peptide; Direct protein sequencing.
SEQUENCE 5 AA; 570 MW; 71A9C9C862A00000 CRC64;
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                                                             0; Indels
   Length
44.4%; Score 12; DB 1; Le
100.0%; Pred. No. 1.6e+06;
ive 0; Mismatches 0;
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P83568;
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Best Local S
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MEDLINE=98163340, PubMed=9504812;
Dunn M.J., Corbett J.M., Wheeler C.H.;
J. Corbett J.M., Wheeler C.H.;
"HSC-2DPAGE and the two-dimensional gel electrophoresis database of dog heart proteins ";
Electrophoresis 18:2795-2802(1997),
-: CATALYTIC ACTIVITY: D-glyceraldehyde 3-phosphate = glycerone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR00065; Triophos ismrse.
PROSITE; PS00171; TIM; PARTIAL.
Direct protein sequencing; Fatty acid biosynthesis; Gluconeogenesis; Glycolysis; Isomerase; Pentose shunt.
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                                                                                                                                                                                                                                                                                                                                                                                                                             Canis Familiaris (Dog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Carnivora; Fissipedia; Canidae; Canis.
NCBI_TaxID=9615;
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SEQUENCE.
MEDLINE=92270459; PubMed=1821096; DOI=10.1016/0196-9781(91)90190-Z;
                                                                                                                                 ;
                                                                                                                                                                                                                                                                                                                                      01-0CT-1996 (Rel. 34, Last Sequence update)
25-0CT-2004 (Rel. 45, Last annotation update)
Triosephosphate isomerase (EC 5.3.1.1) (TIM) (Triose-phosphate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
05-JUL-2004 (Rel. 44, Last sequence update)
Anthor-Klamide I (Contains: AnthorRamide II).
Anthopleura elegantissina (Sea anemone)
Eukaryota; Metazoa; Cnidaria; Anthozoa; Zoantharia; Actiniaria;
Nynantheae; Actiniidae; Anthopleura.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .;
0
                                                                            44.4%; Score 12; DB 1; Length 5; 100.0%; Pred. No. 1.6e+06; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               i; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 5;
  1 1
5 AA; 654 MW; 72CB19C9C0300000 CRC64;
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Pred. No. 1.6e+06;
                                                                                                                                                                                                                                                                                                5 A.A.
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                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                        STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                 isomerase) (Fragment).
                                                                                          Local Similarity
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01-OCT-1996
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ID TPIS_CANPA

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FYRI ANTEL
AC PS 9706,
DT 28-FFB-
DT 05-JUL-
DB Antho-F
OS Anthop)
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CC Nymanth
OX NCB1 TR
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Nothacker H.-P., Rinehart K.L. Jr., McFarlane I.D., Grimmelikhuijzen C.J.P., "Isolation of two novel neuropeptides from sea anemones: the unusual, biologically active L.3-phenyllactyl-Tyr-Arg-Ile-NH2 and its despeptides 12:1165-1173 (1991).
                                                                                                                                                                                                                                                                                                                                                                                                             C.J.P.;
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MEDLINE=22197108; PubMed=12207899; DOI=10.1016/S0006-291X(02)02036-3;
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MEDILINE-294,
Zatylny C., Gagnon J., Boucaud-Camou E., Henry J.;
"ILME: a waterborne pheromonal peptide released by the eggs of Sepia officinalis.";
Biochem. Biophys. Res. Commun. 275:217-222 (2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bukaryota; Metazoa; Mollusca; Cephalopoda; Coleoidea; Neocoleoidea;
Decapodiformes; Sepioidea; Sepiidae; Sepia.
NCBI_TaxID=6610;
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"Fertilization in Sepia officinalis: the first mollusk sperm-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE, SUBCELLULAR LOCATION, TISSUE SPECIFICITY, AND MASS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 4;
1.6e+06;
ches 0; Indels
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100.0%; Pred. No. 1.6e+06;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Direct protein sequencing; Pheromone.
SEQUENCE 4 AA; 505 MW; 6B1697203000000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-MAR-2004 (Rel. 43, Created)
29-MAR-2004 (Rel. 43, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Pheromone peptide ILME.
Sepia officinalis (Common cuttlefish).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13.3%; Score 9; DB 1
50.0%; Pred. No. 1.6e
Conservative 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4 AA
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Matches 2; Conservative
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P01157;
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GRWM HUMAN
Matches
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MEDLINE=55009901:
MEDLINE=55009901:
Merrick B.A., Patterson R.M., Wichter L.L., He C., Selkirk J.K.;
"Separation and sequencing of familiar and novel murine proteins using preparative two-dimensistonal gel electrophoresis.";
Electrophoresis 15:735-745 (1994).
-!- MISCELLANEOUS: On the 2D-gel the determined pI of this unknown protein is: 6.6, its MW is: 19 kDa.

Direct protein sequencing.
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Medicinal lobe;
MEDLIULAR LOCATION:
MEQUELIULAR LOCATION:
MEQUELIULAR LOCATION:
MEQUELIULAR LOCATION:
MEDLIULAR LOCATION
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Eukaryota; Metazoa; Mollusca; Cephalopoda; Coleoidea; Neocoleoidea;
Decapodiformes; Sepioidea; Sepiidae; Sepia.
                                                                                                                                                                                                                                                                                                                                                     01-0CT-1994 (Rel. 30, Created)
1-0CT-1994 (Rel. 30, Last sequence update)
05-JUD-2004 (Rel. 34, Last annotation update)
05-JUD-2004 (Rel. 44, Last annotation update)
UNKNOWN protein from D-PAGE of fibroblasts (P19) (Fragment).
Buks musculus (Mouse).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Musinae; Muscul TaxID=10090;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE, FUNCTION, MASS SPECTROMETRY, AND AMIDATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Amidation; Direct protein sequencing; Neuropeptide. MOD_RES 2 Tryptophan amide. SEQUENCE 2 AA; 261 MW; 737810000000000 CRC64;
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29-MAR-2004 (Rel. 43, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
                                                                                                                                                                                                                                                                                             5 AA.
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P83570;
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GWA_SEPOF
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22.2%; Score 6; DB 1; Length 2; 100.0%; Pred. No. 1.6e+06;

Best Local Similarity

Query Match

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Schleäinger D.H., Pickart L., Thaler M.M.;
Schleäinger D.H., pickart L., Thaler M.M.;
"Growth-modulating serum tripeptide is glycyl-histidyl-lysine.";
Experientia 33:324-325(1977).
-!- MISCELLANEOUS; This serum tripeptide has been found to stimulate growth of some cell types and to inhibit other types in vitro.
GO; GO:0001558; P:regulation of cell growth; NAS.
Gaps
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
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Indels
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SEQUENCE 3 AA; 340 MW; 6331E8100000000 CRC64;
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                                                                                                                                                                                                                                                21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Homo sapiens (Human)
                                                                                                                                                                                                          3 AA.
0; Mismatches
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                                                                                                                                                                                                          PRT;
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Best Local Similarity 100...
1; Conservative
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version	- 2005
GenCore	(c) 1993
	Copyright

OM protein - protein search, using sw model

March 23, 2005, 15:19:16; Search time 68 Seconds (without alignments) 28.438 Million cell updates/sec Run on:

27 l ffglm 5 Perfect score: Sequence:

BLOSUM62 Scoring table:

Gapop 10.0 , Gapext 0.5

2105692 seqs, 386760381 residues Searched:

Total number of hits satisfying chosen parameters:

45841

seq length: 0 seq length: 5 Minimum DB Maximum DB Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

A_Geneseq_16Dec04:* 1: geneseqp1980s:*

geneseqD1990s:* geneseqD2000s:* geneseqD2001s:* geneseqp2002s:* geneseqp2003as:* geneseqp2003bs:*

geneseqp20048:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Result No.	Score	Query Match	Length	DB	ID	Description
	27	100.0	S	7	AAR33009	Aar33009 Alpha-sub
7	27	100.0	5	~	AAR33008	Aar33008 Alpha-sub
e	27	100.0	S	~	AAR33007	Aar33007 Alpha-sub
4	27	100.0	Ŋ	7	AAR33010	Aar33010 Alpha-sub
S	27	100.0	S	7	AAR54549	Aar54549 Cholecyst
9	27	100.0	5	~	AAR54551	Aar54551 Cholecyst
7	27	100.0	S	7	AAR54550	Aar54550 Cholecyst
۵	27	100.0	S	7	AAR54548	Aar54548 Cholecyst
6	27	100.0	S	~	AAW41687	Aaw41687 Tetrapept
10	27	100.0	Ŋ	7	AAW99643	Aaw99643 Substance
11	27	100.0	'n	~	AAY50325	Aay50325 Neutrophi
12	27	100.0	ហ	~	AAW92660	Aaw92660 Human tac
13	27	100.0	Ŋ	ო	AAB23025	Aab23025 Human/rat
14	27	100.0	ഗ	ო	AAY67576	Aay67576 P antagon
15	27	100.0	S	4	AAB91428	Aab91428 Tachykini
16	27	100.0	S	Ŋ	ABB10088	Abb10088 Substance
17	27	100.0	S	ស	AAU77845	
18	27	100.0	ហ	7	ADE94203	Ade94203 High acti
19	27	100.0	S)	7	ADF92530	Adf92530 Substance
20	27	100.0	2	ω	ADN95078	Adn95078 Mammalian
21	27	100.0	Ŋ	œ	ADR43771	Adr43771 Human mag
22	24	88.9	S	~	AAW92702	Aaw92702 Human tac
23	24	ω.	2	ഗ	ABB10089	Abb10089 Substance
24	24	88.9	2	7	ADE94204	Ade94204 High acti
25	22	81.5	ß	~	AAR27697	Aar27697 Cyclic ta

	Aar47524 GHRP-6 an Aaw13221 Growth ho
5 2 AAW92703 5 2 AAW92701 4 2 AAW92701 4 3 AAW31075 4 3 AAW23026 4 4 AAW31075 4 4 AAW37091 4 5 AAW17846 4 7 ADE94198 4 7 ADE94198 5 4 AAW9372 5 5 AAW80389 5 5 AAW80389 5 6 AAW80389 5 7 ADE94205 5 8 AAW80134 5 2 AAW80134	5 2 AAR47524 5 2 AAW13221
	20 74.1 20 74.1
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ALIGNMENTS

RESULT 1

AAR33009 standard; peptide; 5 AA. (revised)
(first entry) 25-MAR-2003 02-APR-1993 AAR33009; AAR33009

Alpha-substituted short peptide.

CCK; neuropeptide; endorphin; hormone; LHRH; contraception; analgesia; improved bioavailability. Location/Qualifiers Key Modified-site Synthetic.

/note= "alpha-Me-Leu"

/note= "Met-NH2" WO9219254-A1

Modified-site

92WO-US003119. 15-APR-1992; 12-NOV-1992.

(WARN) WARNER LAMBERT CO.

91US-00690755. 92US-00852086.

24-APR-1991; 20-MAR-1992;

Howson W; Horwell DC, Hughes J, Richardson RS,

WPI; 1992-398522/48.

New alpha-substd. polypeptide are e.g. selective receptor ligands - for treating inflammation, pain, stroke, ulcers, hypertension, heart failure, depression, cancer, asthma, psychosis, arthritis, etc.

Claim 3; Page 41; 46pp; English.

The peptide is a specifically claimed example of a group of generically claimed mono-, di-, tri-, tetra- and penta-peptides which include a substituent on an alpha-C atom in the chain. Such substitution may modify the bioavailability, stability, or absorbability of the peptide and hence may improve the activity of the peptide as a drug. Depending on the

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nature of the parent peptide (hormone, endorphin, CCK, NK2, chemotactic peptide, etc.), the modified peptides are variously useful for treating obesity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, addictive drug withdrawal symptoms, hypertension, heart failure, asthma, bladder dysfunction, psychosis and arthritis; and as contraceptives. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PN field.) (Updated on field.) (Updated on 25-MAR-2003 to correct PN field.)
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                                                                                                                                                                             100.0%; Score 27; DB 2; Length 5; 100.0%; Pred. No. 1.8e+06; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "alpha-Me-Phe"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                AAR33008 standard; peptide; 5 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                Alpha-substituted short peptide.
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                                                                                                                                                          Query Match
Query Match
Best Local Similarity 100.0*
S; Conservative
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92US-00852086
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(first entry)
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                                                                                                                                             Sequence 5 AA;
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Modified-site
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02-APR-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
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cognition or memory disorders, spasticity, depression, diabetes, cancer, asthma, bladder dysfunction, psychosis and arthritis; and as contraceptives. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PR field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New alpha-substd. polypeptide are e.g. selective receptor ligands - for treating inflammation, pain, stroke, ulcers, hypertension, heart failure, depression, cancer, asthma, psychosis, arthritis, etc.
                                                                                                                                                                                                                                                                                                                                                                                       CCK; neuropeptide; endorphin; hormone; LHRH; contraception; analgesia;
improved bioavailability.
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                                                                                                              100.0%; Score 27; DB 2; Length 5; 100.0%; Pred. No. 1.88+06; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "alpha-Me-Phe"
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                              AAR33007 standard; peptide; 5 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "Met-NH2"
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92US-00852086.
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(first entry)
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                                                                                                         Query Match
Best Local Similarity
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02-APR-1993
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Matches
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The sequences given in AAR53117-38 and AAR54530-51 are peptide analogues of cholecystokinin (CCK) which can be used to treat obesity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, hypertension, heart failure, cognition, memory enhancement, spasticity, depression, diabetes, cancers, asthma, bladder dysfunction, psychosis, arthritis and in the treatment of substance withdrawl. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                   Peptide analogue; peptoid; cholecystokinin; CCK; obesity; anxiety; gastrointestinal ulcers; pain; stroke; inflammation; hypertension; heart failure; cognition; memory enhancement; spasticity; depression; diabetes; cancers; asthma; bladder dysfunction; psychosis; arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New cholecystokinin analogues - useful e.g. in treatment of pain, obesity, stroke, anxiety, and gastrointestinal ulcers.
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0; Mismatches
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                                                                                                                                                                  AAR54549 standard; peptide; 5 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /label= MePhe
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93US-00131693.
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(first entry)
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14-DEC-1994
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Matches
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AAR54551
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                                                                       100.0%; Score 27; DB 2; Length 5; 100.0%; Pred. No. 1.8e+06;
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field.) (Updated on 25-MAR-2003 to correct PI field.)
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Pred. No. 1.8e+06;
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100.0%;
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                                                                                                            5; Conservative
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improved bioavailability
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02-APR-1993
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14-OCT-1993;
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14-DEC-1994
                             Synthetic
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AAR54548
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                                                                     Peptide analogue, peptoid; cholecystokinin, CCK; obesity, anxiety; gastrointestinal ulcers; pain; stroke; inflammation; hypertension; heart failure; cognition; memory enhancement; spasticity; depression; diabetes; cancers; asthma; bladder dysfunction; psychosis; arthritis.
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                                                                                                                                                                                                                                                                                                                    New cholecystokinin analogues - useful e.g. in treatment of pain, obesity, stroke, anxiety, and gastrointestinal ulcers.
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                                                                                                                                                                 /note= "Amidated C-terminal"
                                                    Cholecystokinin analogue peptide #44.
                                                                                                                                        Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AARS4550 standard; peptide; 5 AA.
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                                                                                                                                                       label= MeMet
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93US-00131693
                             (revised)
(first entry)
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5; Conservative
                                                                                                                                                                                                                                                                                  Horwell DC, Howson W,
                                                                                                                                                                                                                                                                                                     WPI; 1994-151243/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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                                                                                                                                     Key
Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 5 AA;
                                                                                                                                                                                                                    14-OCT-1993;
                           25-MAR-2003
                                    14-DEC-1994
                                                                                                                                                                                  WO9409031-A1
                                                                                                                                                                                                                                      19-OCT-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
14-DEC-1994
                                                                                                                    Synthetic
          AAR54551;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR54550;
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The sequences given in AARS3117-38 and AARS4530-51 are peptide analogues of cholevstokinin (CCK) which can be used to treat obesity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, hypertension, heart failure, cognition, memory enhancement, spasticity, depression, diabetes, cancers, asthma, bladder dysfunction, psychosis, arthritis and in the treatment of substance withdrawl. (Updated on 25-MAR-2003 to correct PN
cancers; asthma; bladder dysfunction; psychosis; arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Peptide analogue, peptoid, cholecystokinin, CCK, obesity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, hypertension, heart failure, cognition, memory enhancement, spasticity, depression, diabetes, cancers, asthma, bladder dysfunction, psychosis, arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New cholecystokinin analogues - useful e.g. in treatment of pain, obesity, stroke, anxiety, and gastrointestinal ulcers.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 27; DB 2; Length 5; 100.0%; Pred. No. 1.8e+06; rive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Horwell DC, Howson W, Hugues J, Richardson RS;
                                                                                                                                                                                                                       /note= "Amidated C-terminal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "Amidated C-terminal"
                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cholecystokinin analogue peptide #41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 3; Page 66; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR54548 standard; peptide; 5 AA.
                                                                                                                                                                   /label= MeLeu
                                                                                                                                                                                                                                                                                                                                                                                        93WO-US009809
                                                                                                                                                                                                                                                                                                                                                                                                                                        92US-00963169.
93US-00131693.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (WARN ) WARNER LAMBERT CO.
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Best Local Similarity
Matches 5; Consery
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corneal disorder remedy for corneal ulcer, corneal epithelial peeling,
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                                                                                                                                                   Sequence 5 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
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Best Local
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                                                                                                                                                                                                         The sequences given in AAR53117-38 and AAR54530-51 are peptide analogues of cholecystokinin (CCK) which can be used to treat obesity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, hypertension, heart failure, cognition, memory enhancement, spasticity, depression, diabetes, cancers, asthma, bladder dysfunction, psychosis, arthritis and in the treatment of substance withdrawl. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ophthalmic drug; corneal disorder; ulcer; epithelial peeling; dry eye; keratitis; insulin like growth factor-1; IGF-1; eye drop.
                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ophthalmic drug composition containing tetra:peptide - is useful as
                                                                                                                                                            New cholecystokinin analogues - useful e.g. in treatment of pain, obesity, stroke, anxiety, and gastrointestinal ulcers.
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                                                                                                                                                                                                                                                                                                                        100.0%; Score 27; DB 2; I 100.0%; Pred. No. 1.8e+06; iive 0; Mismatches 0;
                                                                                                                   Hugues J, Richardson RS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "C-terminal amide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW41687 standard; peptide; 5 AA.
                                                                                                                                                                                           Claim 3; Page 66; 73pp; English
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                                          93WO-US009809
                                                              92US-00963169
93US-00131693
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                                                                                                                                                                                                                                                                                                            Ouery Match
Best Local Similarity 100.vv
                                                                                             (WARN ) WARNER LAMBERT CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-JUN-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nishida T, Nakamura M,
                                                                                                                  Horwell DC, Howson W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1998-076907/07.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tetrapeptide #4
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                                          14-OCT-1993;
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                                                              19-OCT-1992;
                                                                        08-OCT-1993;
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                     28-APR-1994
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                                                                                                                                                                                                                                                                                                                                                                   7
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                                                            ophthalmic drug composition which contains phe-Gly-Leu-Met-NH2 or its medicinally acceptable salts as the active ingredient. It is used, together with insulin like growth factor-I (IGF-I), to treat corneal disorders such as corneal ulcer, corneal epithelial peeling, dry eye and keratitis. The dosage is 0.1-5000 (preferably 1-1000) mg/day of the active ingredient and 0.001-100 (preferably 0.01-10) mg/day of IGF-I. The preferable form of the composition is eye drops
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A composition has been developed for supplying a peptide to an opioid receptor or that interferes with binding of substance P to its receptor. The composition comprises: (a) myogenic cells that contain heterologous DNA encoding the peptide to express the peptide; and (b) a pharmaceutically acceptable carrier. The composition is useful for relieving pain and for treating behavioural and perceptive abnormalities using myoblast transfer therapy. It is useful in a method for treating psychiatric conditions that involve abnormal perception e.g. depression, chornic anxiety syndromes, paranois, alcoholism and drug addiction, chronic pain and other diseases in which opioid neurons and substance P sensitive neurons play a role. The composition provides a continuous,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New composition for supplying peptide to opioid receptor - comprises myogenic cells containing heterologous DNA encoding peptide and carrier.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Substance P; myoblast transfer therapy; pain relief; analgesic; behavioural abnormality; perceptive abnormality; opioid receptor; psychiatric condition; depression; chronic anxiety syndrome; paranoia; alcoholism; drug addiction; chronic pain; neuron.
                                        sequence is shown in the specification. The invention relates to
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                                                                                                                                                                                                                                                                                             100.0%; Score 27; DB 2; Length 5; 100.0%; Pred. No. 1.8e+06; rive 0; Mismatches 0; Indel8
                                                                                                                                                                                                                                                                                                  Length 5;
Disclosure; Page 11; 19pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW99643 standard; peptide; 5 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Substance P analogue peptide.
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                                                                                                                                                                                                                                                                                                                        Similarity
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This invention describes a novel method for the use and preparation of cell activating compositions which involves preparing a cell activating compositions which involves preparing a cell activating about neutral or higher pH to produce a homogenate; (b) removing cabout neutral or higher pH to produce a homogenate; (b) removing the homogenate in the homogenate in the a protease; and (d) carticulates from the homogenate and selecting fractions that exhibit cell cartivation activity. The methods can be used for improving treatment controme or reducing risk of treatment of e.g. cardiovascular disease, cutoment of inflammatory disease, trauma, autoimmune diseases, arthritis, organ controme or reducing risk of treatment of e.g. cardiovascular disease, contromed and diabetic complications, stroke, ischemia, creptorion, diabetes and diabetic complications, stroke, ischemia, cretinopathy, diabetes, venous insufficiency, unstable angina or trauma. They can be used in the veterinary treatment of a non-human subject. They can be used in the used to lower cell activation resulting from these diseases and deficiencies. The detection of an elevated level of hydrogen peroxide can be used to detect an inflammatory condition. An
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long term supply of opioid peptides (long-term analgesia) which lasts for up to at least 6 years. The present sequence represents a specifically claimed substance P analogue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cell activation; pancreas; treatment; cardiovascular disease; trauma; inflammatory disease; autoimmune disease; arthritis; diabetes; stroke; organ rejection; isohemia; Alzheimer's disease; myocardial infarction; haemorrhagic shock; diabetic retinopathy; venous insufficiency; angina; trauma; protease inhibitor; hypertension; sepsis.
                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                           Neutrophil-activating pancreatic derived peptide 125.
                                                                                                                 100.0%; Score 27; DB 2; I 100.0%; Pred. No. 1.8e+06;
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                                                                                                                                                        0; Mismatches
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                                                                                                                                                                                                                                                                                                               AAY50325 standard; peptide; 5 AA,
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                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              activation;
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Best Local S
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elevated level of hydrogen peroxide in plasma or whole blood and in the presence of superoxide dismutase (SOD) indicates leukcyte up regulation, as disease onset of the onset of an acute cardiovascular disorders, such peroxide in plasma or whole blood and a low level in the presence of SOD hypertension or sepsis. AMYSO201-YSO334 represent peptides used in the manner of SOD method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes a method for screening compounds for inhibiting a neurcroxin. The method involves incubating tachykinin agonists with neuronal cells and a beta-amyloid peptide neurotoxin. The methods can be used for identifying compounds for treating diseases characterised by an Down's syndrome, and the syndromes of hereditary cerebral haemorrhage with amyloidosis and non-inherited congophilic angiopathy with cerebral beta-amyloid peptide fragments. (Updated on 20-MAR-2003 to correct PF leield.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tachykinin agonist; beta-amyloid; inhibition; neurotoxin; treatment; Alzheimer's disease; Down's syndrome; amyloidosis; human; hereditary cerebral haemorrhage; non-inherited congophilic angiopathy.
                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Screening for neurotoxin inhibitors - by testing compounds for their effect on beta-amyloid peptide neurotoxic effect on neuronal calls.
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                                                                                                                                                                   100.0%; Score 27; DB 2; Length 5; 100.0%; Pred. No. 1.8e+06; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                     Human tachykinin agonist beta-amyloid peptide fragment #6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure, Col 13-14; 28pp; English.
                                                                                                                                                                                                                                                                                                                                AAW92660 standard; peptide; 5 AA.
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(first entry)
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Best Local Similarity
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30-APR-1999
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Matches
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Length 5;

Score 27; DB 2; Length 5; Pred. No. 1.8e+06; Mismatches 0; Indels

/ Match 100.0%; Sc Local Similarity 100.0%; Pi hes 5; Conservative 0;

Query Match

Matches

(first entry)

19-MAY-2000

AAY67576;

P antagonist peptide #4.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The inevntion relates to a method for detecting magnesium binding defects. The method comprises quantitating a tachykinin C-terminal pentapeptide (e.g., AAB23025) and its degradation products (e.g., AAB23026) in blood using an antibody specific for the generalised mammalian tachykinin C-terminal pentapeptide Phe-(Phe/Val)-Gly-Leu-Met-NH2 (AAB23028). The method is useful for detecting cellular magnesium binding defects which are associated with abnormal physiological states such as sodium-sensitive essential hypertension and type 2 diabetes mellitus. The present sequence represents the C-terminal 5 amino acids of the tachykinin Substance P (AAB23027) from human and rat, which may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    physiological states such as sodium-sensitive essential hypertension and type 2 insulin-resistant diabetes mellitus, comprises measuring a specific pentapeptide in blood.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                               Substance P; tachykinin; human; rat; magnesium binding defect; sodium sensitive essential hypertension; insulin resistance; type 2 diabetes; antibody; immunoassay; quantification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detecting magnesium binding defects associated with abnormal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 27; DB 3; Length 5; ilarity 100.0%; Pred. No. 1.8e+06; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                 Human/rat tachykinin Substance P C-terminal pentapeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    assayed according to the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "C-terminal amide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
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                                                                                                                                                                 AAB23025 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-MAR-2000; 2000WO-US003707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-00265690
                                                                                                                                                                                                                                                                 (first entry
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Best Local Similarity
Matches 5; Conserv
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                              PFGLM
FFGLM
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                                                                                                                                                                                                                                                                    16-JAN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rattus sp
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                                                                                                                                                                                                                    AAB23025;
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JD AAB23025
AAC AAB2
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The invention provides a pharmaceutical and/or veterinary formulation that comprises 2 -30% of active agents which include a gonadotropin-releasing hormone (GnRH) agonist, 0.5 - 20% of a pore-forming agent which is not lecithin, and the remainder stearin. The formulation is useful as a sustained release implant which can deliver the active agent for a period of 7 days to 2 years. Sequences AAY67573-578 represent P antagonist peptides used in the composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bioimplant formulation for sustained delivery of an active agent over 7 days to 2 years, comprises active agent, pore-forming agent and stearin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Protection, endogenous therapeutic peptide; peptidase; conjugation, blood component, modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter.
Pharmaceutical, veterinary, gonadotropin-releasing hormone, GRRH, pore-forming agent, lecithin, stearin, P antagonist.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 27; DB 3; Length 5; 100.0%; Pred. No. 1.8e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                     /note= "C-terminal amide"
                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rachykinins peptide SEQ ID NO:604.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB91428 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rathjen DA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 20; Page 21; 37pp; English.
                                                                                                                                                                                                                                                                                99WO-AU000585.
                                                                                                                                                                                                                                                                                                                          98AU-00004730.
                                                                                                                                                                                                                                                                                                                                             98AU-00004731.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Walsh JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-182528/16.
                                                                                                                                                                                                                                                                                                                                                                                                             (PEPT-) PEPTECH LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
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                                                                                                                                                                                               WO200004897-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 5 AA;
                                                                                                                               Modified-site
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Synthetic.
                                                                    Inidentified
                                                                                                                                                                                                                                                                              20-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                20-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                   13-MAY-1999;
                                                                                                                                                                                                                                      03-FEB-2000
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AAB91428
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AAY67576 standard; peptide; 5 AA.

RESULT 14 AAY67576 ID AAY67

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1 FFGLM

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WO200069900-A2

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comprising a therapeutically active amino acid region (III) and a reactive group (II) (e.g. succinimidyl and maleimido groups) attached to a less therapeutically active amino acid region (IV), which covalently conds with amino/hydroxyl/thiol groups on blood components to form a peptidase stabilised therapeutic peptide composed of 3.50 amino acids. (I) are useful for modifying therapeutic peptides e.g. hormones, growth vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent work for modifying and attaching therapeutic peptides to allowin prevents or reduces the action of peptidases to increase length of activity (half intracellular uptake and interference with physiological processes. ABB90829 to AAB9241 represent invention
                                                                                                                                                                                                                  Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     healing; insulin-like growth factor-I; tear; abrasion; skin ulcer;
                                                                                                                                                                                                                                                                                     The present invention describes a modified therapeutic peptide (I)
                                                                                                                                                                      Thibaudeau K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 27; DB 4; Length 5; 100.0%; Pred. No. 1.8e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Substance P analog used in wound healing treatment#11.
                                                                                                                                                                    Holmes DL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                          Disclosure; Page 397; 733pp; English
                                                                                                                                                                    Milner PG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABB10088 standard; peptide; 5 AA.
                                                    17-MAY-2000; 2000WO-US013576.
                                                                                              99US-0153406P.
                                                                                    99US-0134406P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-AUG-2001; 2001WO-JP006933
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-AUG-2000; 2000JP-00242489.
28-NOV-2000; 2000JP-00361388.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   surgical incision; burn.
                                                                                                                                                                Bridon DP, Ezrin AM,
                                                                                                                                     (CONJ-) CONJUCHEM INC.
                                                                                                                                                                                           WPI; 2001-112059/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 5 AA;
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                                                                                17-MAY-1999;
10-SEP-1999;
                                                                                                            15-OCT-1999;
                            23-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABB10088;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 16
ABB10088
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                                                                                                                                                                                            The invention relates to skin wound healing promoters, containing substance P analogs or their pharmaceutically-acceptable salts, and insulin-like growth factor-I as the active ingredient. The promoters are for treating wounds like tears, abrasions, surgical incisions, or skin ulcers and burns. The current sequence represents a substance P analog for use in wound healing treatment
                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tachykinin; substance P; hypertension; hypotensive; antidiabetic; gynaecological; salt-insensitive hypertension; magnesium binding; insulin resistance; type 2 diabetes mellitus; pre-eclampsia; eclampsia.
                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                     Skin wound healing promoters or skin epidermal extension promoters containing substance P analogs and insulin-like growth factor-I for treating wounds like tear, abrasion, surgical incision, skin ulcers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             are
e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New monopeptides derived from butadienes, ethylenes and propanes; magnesium binding defect antagonists, useful in the treatment of hypertension, insulin resistance of type 2 diabetes mellitus and
                                                                                                                                                                                                                                                                                                                                                    ;
0
                                                                                                                                                                                                                                                                                                                    100.0%; Score 27; DB 5; Length 5; 100.0%; Pred. No. 1.8e+06; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                              0; Indels
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/note= "C terminal-amide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tachykinin N -terminal pentapeptide.
                                                                                                                                                                     Disclosure; Page 4; 20pp; Japanese.
                                                       Nakamura M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAU77845 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (MAGN-) MAGNESIUM DIAGNOSTICS INC.
             (SANT ) SANTEN PHARM CO LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-AUG-2001; 2001WO-US024909.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-AUG-2000; 2000US-00635266.
                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.00
Then 5; Conservative
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                                                   Nishida T, Nakata K,
                                                                             WPI; 2002-269153/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-280663/32.
                          (NISH/) NISHIDA T.
                                                                                                                                                                                                                                                                                                                                                                      1 FFGLM 5
                                                                                                                                                                                                                                                                                                                                                                                               1 FFGLM 5
                                                                                                                                                                                                                                                                                       Sequence 5 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-FEB-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAU77845;
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                                                                                                                                               burns.
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Disclosure; Page 2; 38pp; English.

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This invention relates to novel therapeutic compounds and methods used for treating mammals with disorders such as salt-insensitive bypertension. The monopeptide compounds of the invention are derived from butadienes, ethylenes and propanes. The compounds of the invention are used to correct a defect in magnesium binding within the plasma membranes of somatic cells which results in a decrease in the intracellular concentration of magnesium ions. These compounds may be used in the treatment of a mammal affected with magnesium binding defect, saltsensitive (particularly hypertension), insulin resistance of type 2 diabetes mellitus and pre-eclampsia/eclampsia. The compounds of the invention have an advantage over prior art compounds in that these compounds are biologically stable. The present sequence represents the a pentapeptide from the C terminal sequence of tachykinin known as a butter of a sufficient to correct the magnesium binding defect responsible for causing hypertension \$\$666666666666666666888

Sequence 5 AA;

Length 5; Indels 100.0%; Score 27; DB 5; L 100.0%; Pred. No. 1.8e+06; Live 0; Mismatches 0; Conservative Local Similarity FFGLM 5 Query Match Matches ò 셤

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> 'n FFGLM

ADE94203 standard; peptide; 5 AA. ADE94203; ADE94203

LD ADE9

AC ADE9

XX ADE9

(first entry) 12-FEB-2004 High activity minimal IGF-1-derived peptide fragment #15.

ophthalmological; dermatological; vulnerary; insulin growth factor 1; IGF-1; ophthalmology; dermatology; keratic injury; wound healing; skin; corneal ulcer; exofoliation of corneal epithelium; keratitis; dry eye; scratch; surgical cutting; skin ulcer; burns.

Homo sapiens

/note= "amidated C-terminus" Location/Qualifiers 5 Misc-difference

WO2003048192-A1

12-JUN-2003.

03-DEC-2002; 2002WO-JP012632

03-DEC-2001; 2001JP-00368103

(SANT) SANTEN PHARM CO (NISH/) NISHIDA T.

Nakamura M; Inui M, Nishida T,

WPI; 2003-505280/47.

Novel peptides based on minimum activity expression units of insulin-like growth factor-1, applicable in remedies in ophthalmology and dermatology for treating keratic injury and promoting wound healing in skin. Novel peptides based growth factor-1, appl

Disclosure; Page 7; 25pp; Japanese.

The invention relates to the determination of the smallest peptide fragment of insulin growth factor 1 (IGF-1) with the highest activity for use in ophthalmology and dermatology. The peptides are applicable in

ö The invention describes a chimeric hybrid molecule (I) of a cyclic alkaloid moiety which binds as an agonist to a mammalian/human mu opioid receptor and a peptide moiety which binds as an agonist to a mammalian/human substance P. (I) is useful for inhibiting development of opioid tolerance by chemically combining a pharmacologically active form of substance P with morphine in (I). (I) is useful for transporting an active form of S P or neuropepiide across the blood-brain barrier into the central nervous system using the active metabolite of morphine, morphine of reciprocally regulating analgesic chemicals across the blood-brain barrier into the archive manalagesic chemicals across the blood-brain barrier into the central nervous system using (I). (I) is useful for analgesic; Mu-opioid receptor agonist; substance P receptor agonist; chimeric hybrid; cyclic alkaloid moiety; mu opioid receptor; substance P; opioid tolerance; morphine; substance P; SP; neuropeptide; blood-brain barrier; morphine 6-glucuronide; pain; drug abuse; analgesia; tolerance development; dependence formation; Chimeric hybrid molecule useful for treating pain comprising cyclic alkaloid molety which binds as agonist to mammalian to human mu opioid receptor and peptide molety which binds as agonist to mammalian substance remedies in ophthalmology and dermatology for treating keratic injury and promoting wound healing in the skin. The keratic injury is particularly corneal uluer, exofoliation of corneal epithelium, keratitis or dry eye. The skin wound can be scratches, surgical cutting, skin ulcer, or burns. This sequence represents one of the peptides of the invention with IGF-1 Gaps ö Length 5; 0; Indels ; DB 7; L Score 27; DB Pred. No. 1.86; Mismatches /note= "C-terminal amide" Location/Qualifiers ADF92530 standard; peptide; 5 AA. Substance P receptor agonist #3. ő 7; Page 7; 11pp; English. 100.0%; 26-APR-2002; 2002US-00134187. substance P receptor agonist 26-APR-2002; 2002US-00134187 (first entry) 5; Conservative WPI; 2003-900618/82 Similarity Ŋ S US2003202981-A1. ĸ FFGLM Sequence 5 AA; Key Modified-site (KREA/) KREAM Unidentified 26-FEB-2004 30-OCT-2003 Query Match Best Local S Н ADF92530; Kream RM; activity Claim Matches RESULT 19 ADF92530 8888888888 ઠે 셤

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treating pain in a mammal and for treating drug abuse in a mammal by administering (I) in substitution for the drug on which the mammal became dependent and/or tolerant and thereafter adjusting the dosage as tolerance and/or dependence is modulated. (I) induces analgesia in a mammal with tolerance development markedly less than that of morphine. (I) efficiently modulates the activation of the MOR and to reduce or eliminate tolerance development and dependence formation. This is the amino acid sequence of a peptide that functions as a substance P receptor
                                                                                                                                                                                                                                                                                               Sequence 5 AA;
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100.0%; Score 27; DB 7; Length 5; ilarity 100.0%; Pred. No. 1.8e+06; Conservative 0; Mismatches 0; Indels Query Match Best Local Similarity Matches 5; Conserv Ŋ 1 FFGLM ò

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1 FFGLM

RESULT 20

ADN95078 standard; peptide; 5 AA ADN95078;

(first entry) 26-AUG-2004

Mammalian substance P peptide (amino acids 7-11).

Opioid tolerance; substance P; morphine; cyclic alkaloid; mammalian; mu opioid receptor; acute pain; chronic pain; drug abuse; opioid analgesia; analgesic; antiaddictive.

Mammalia.

note= "C-terminal amide" Location/Qualifiers Key Modified-site

US2004106636-A1

03-JUN-2004.

17-OCT-2003; 2003US-00688741.

26-APR-2002; 2002US-00134187

(KREA/) KREAM R M.

Kream RM;

WPI; 2004-419489/39.

Inhibiting development of opioids tolerance involves use of chimeric hybrid molecules containing an opioid moiety of chemically modified

Disclosure; SEQ ID NO 3; 10pp; English.

The present invention relates to a method of inhibiting the development combination of an active form of substance P with morphine in a new combination of an active form of substance P with morphine in a new chimeric hybrid molecule. The morphine is chemically modified and covalently linked through its 6'0H group, and comprises a cyclic alkaloid moiety which binds as an agonist to a mammalian or human mu opicid receptor. An active C-mu terminal substance P fragment, chemically peptide and covalently linked through its free NH2 group, comprises a mammalian/human substance P receptor. A compact, but flexible, molecular hinge covalently cross links morphine through its 6'0H group to the free NH2 group of the substance P receptor agonist moiety, so as to allow both

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and the substance P receptor agonist moieties to
                              activate their respective receptors simultaneously and independently. The chimeric hybrid molecules are administered intrathecally, systemically, thradermally, parenterally (e.g. subcutaneously, intravenously), through injection, transdermally, (e.g. topically), transmucosally or pain, and drug abuse. The molecules show reduced side effects. The molecules provide opicid analogesia in living subjects while inhibiting tolerance development and dependence formation. The present sequence representing a peptide from mammalian substance P is used in the method
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Sequence 5 AA;

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ö Length 5; 0; Indels Score 27; DB 8; L Pred. No. 1.8e+06; ; Mismatches 0; . 100.0%; 5; Conservative Query Match Best Local Similarity Matches 5; Conserv

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Gaps

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1 FFGLM

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RESULT 21 ADR4377;

ADR43771 standard; peptide; 5 AA.

ADR43771;

(first entry) 18-NOV-2004

Human magnesium binding defect (MgBD) peptide mimetic #1.

physiological disorder; preeclampsia; pregnancy; salt-sensitive essential hypertension; type 2 diabetes mellitus; human. Magnesium binding defect; MgBD; MgBD binding defect peptide mimetic;

Homo sapiens

Location/Qualifiers Key Modified-site

/label= OTHER /note= "OTHER= C-terminal amide"

US2004171093-A1.

02-SEP-2004.

22-MAR-2004; 2004US-00805881.

10-WAR-1999; 99US-00265690. 09-MUG-2000; 2000US-00635266. 24-JAN-2002; 2002US-0063569. 29-MUG-2002; 2002US-00231333. 28-OCT-2003; 2003US-00695536.

(WELL/) WELLS

Wells IC;

WPI; 2004-625105/60.

Assessing predisposition to physiological disorder associated with magnesium binding defect in individual, by measuring level of amidated peptides associated with magnesium binding defect in sample and comparing peptide level to standard.

Claim 1; SEQ ID NO 1; 21pp; English.

The invention relates to a method of assessing a predisposition to a physiological disorder associated with a magnesium binding defect in an individual, involving measuring the level of amidated peptides associated with the magnesium binding defect in a sample of body fluid of the

cc significantly lower level of the peptide is indicative of a significantly lower level of the peptide is indicative of a control of the individual to the physiological disorder. The individual to the physiological disorder associated with a magnesium binding defect in an individual, involving comparing the level of peptide to the level of peptide is indicative of the progress of treatment of the individual, and the progress of treatment of the individual, and meaning the level of peptide is indicative of the progress of treatment of the individual, and metal of peptide is indicative of the progress of treatment of the individual, and metal of peptide is indicative of the progress of treatment of the individual, and metal of peptide is indicative of the progress of treatment of the individual, and metal of peptide is indicative of the progress of treatment of the individual, and metal of peptide is indicative of the progress of the sembrane tightly bound magnesium in mammalian somatic cells involving the somatic cells from the body fluid by centrifugation, resuspending the somatic cells from the body fluid by centrifugation, resuspending the somatic cells in a cell stabilising buffer, removing a sample of the somatic cells and repeating the removing and measuring steps at subsequent times until the level of tightly bound magnesium is contour to a plasma membrane of mammalian somatic cells having a deficit of plasma membrane tightly bound magnesium in a physiological medium including magnesium in, and and measuring the level of the substance to be tested to the suspension and measuring the level of the sound magnesium and magnesium in the plasma membrane tightly bound magnesium in the plasma membrane tightly bound magnesium of the substance to be tested to the suspension and measuring the level of plasma membrane tightly bound magnesium of the substance to be tested to the substance to be tested is the plasma membrane tightly bound magnesium of the substance to be tested to the substance to be tested to the sub to preeclampsia during pregnancy, salt-sensitive essential hypertension or type 2 diaberes mellitus associated with the magnesium binding defect. The method is also useful for ameliorating or correcting a magnesium binding defect (MgBD) in an individual. This sequence represents a human MgBD mimetic peptide of the invention. ameliorating or correcting a magnessium binding defect in an individual involving administering to the individual a substance which promotes binding of tightly bound magnesium ion to the plasma membrane of mammalian somatic cells. The methods are useful for assessing a predisposition to a physiological disorder associated with a magnesium binding defect in an individual, where the disorder is a predisposition

Sequence 5 AA;

ö Length 5; 0; Indels 100.0%; Score 27; DB 8; I 100.0%; Pred. No. 1.8e+06; Mismatches 0 Query Match 100. Best Local Similarity 100. Matches 5; Conservative

FFGLM 5

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Search completed: March 23, 2005, 15:32:58 Job time : 73 secs

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Run on:

OM protein - protein search, using sw model

March 23, 2005, 14:50:07; Search time 92 Seconds (without alignments) 17.995 Million cell updates/sec

SEQ1 Title: Perfect score:

27 1 ffglm 5 Sequence:

BLOSUM62 Scoring table:

Gapop 10.0 , Gapext 0.5

1407402 seqs, 331100923 residues Searched:

Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 5

Post-processing: Minimum Match 0% Maximum Match 100%

Listing first 45 summaries

Published Applications AA:* Database :

| Cgn2_6/ptodata1/lpubpaa/US07_PUBCOMB.ppp:*
| Cgn2_6/ptodata1/lpubpaa/US07_NEW_PUB.ppp:*
| Cgn2_6/ptodata1/lpubpaa/US06_NEW_PUB.ppp:*
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| Cgn2_6/ptodata1/lpubpaa/US10B_PUBCOMB.ppp:*
| Cgn2_6/ptodata1/lpubpaa/US10D_PUBCOMB.ppp:*
| Cgn2_6/ptodata1/lpubpaa/US0O_NEW_PUB.ppp:*
| Cgn2_6/ptodata1/lpubpaa/US0O_NEW_PUB.ppp:*
| Cgn2_6/ptodata1/lpubpaa/USOO_NEW_PUB.ppp:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result		Query				
No.	Score	Match	Match Length DB	ΩB	ID	Description
	27	100.0	70	6	27 100.0 5 9 US-09-265-690C-1	Sequence 1. Appli
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7	.7	100.0		14	US-10-053-669-1	Sequence 1, Appl
m	27	100.0		15	US-10-134-187-3	Sequence 3, Appl
4	27	27 100.0		16	5 16 US-10-688-741-3	Sequence 3, Appl
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Description	Sequence 1, Appli	Sequence 1, Appli	Sequence 3, Appli	Sequence 3, Appli	Sequence 1, Appli	15,	Sequence 16, Appl	6	Sequence 2, Appli	Sequence 3, Appli	Sequence 2, Appli	Sequence 3, Appli	Sequence 2, Appli	
01	US-09-265-690C-1	US-10-053-669-1	US-10-134-187-3	US-10-688-741-3	US-10-805-881-1	US-10-497-628-15	US-10-497-628-16	US-10-821-240A-270	US-09-265-690C-2	US-10-230-133-3	US-10-053-669-2	US-10-695-536-3	US-10-805-881-2	
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Score	27	27	27	27	27	27	24	22	21	21	21	21	21	
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Sequence 2, Appli Sequence 30, Appl Sequence 17, Appl Sequence 4, Appli	Sequence 4, Appli Sequence 4, Appli Sequence 4, Appli Sequence 24, Appli Sequence 22, Appl	44, AF 44, AF 44, AF 298, 338, A	9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9	3 3 5 7 7 8 8 5 7 9 6 0 9 6 0 9 6 0 9 6 0 9 6 0 9 6 0 9 6 0 9 6 0 9 6 0 9 9 9 9
4440-	0-053-669 0-053-669 0-695-536 0-346-737	4 US-10-155-170-4 4 US-10-351-641-8 6 US-10-822-661-4 1 US-10-821-240A- 1 US-10-920-306-3	US-10-783- US-09-879-4 US-10-620- US-10-620- US-10-337- US-10-891- US-10-891- US-10-891-	3 16 US-10-695-536-2 4 17 US-10-823-964A-11 5 11 US-09-920-306-40 5 14 US-10-301-499A-25 5 14 US-10-168-789A-39 5 15 US-10-194-441A-85 5 15 US-10-131-366-9 5 16 US-10-128-520-360
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ALIGNMENTS

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Sequence 1, Application US/09265690C
Sequence 1, Application US/09265690C
Sequence 1, Application Wo. US20010051345A1
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Method for Detecting Deficient Cellular Membrane Tightly Bound Macritate OF INVENTION: for Disease Diagnosis
TITLE OF INVENTION: for Disease Diagnosis
CURRENT APPLICATION NUMBER: US/09/265,690C
CURRENT FILING DATE: 1999-03-10
STORMARE: PATENTIN VERSION 3.0
SEQ ID NOS: 4
SOFTWARE: PATENTIN VERSION 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      i LOCATION: (5) (5)
COTHER INFORMATION: AMIDATION
US-09-265-690C-1
                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: MOD RES
US-09-265-690C-1
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100.0%; Score 27; DB 9; Length 5; 100.0%; Pred. No. 1.3e+06; ive 0; Mismatches 0; Indels Query Match
Best Local Similarity 100.
Matches 5; Conservative

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Gaps ;

> 1 FFGLM 5 1 FFGLM 5 ò g

RESULT 2 US-10-053-669-1 ; Sequence 1, Application US/10053669 ; Publication No. US20030077658A1

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APPLICANT: Wells, Tibert C.

TITLE OF INVENTION: Method for Detecting Deficient Cellular Membrane Tightly Bound
TITLE OF INVENTION: Magnesium for Disease Diagnosis
TITLE OF INVENTION: Magnesium for Disease Diagnosis
FILE REFERENCE: 800812-0005
CURRENT APPLICATION NUMBER: US/10/805,881
PRIOR APPLICATION NUMBER: US 10/653,669
PRIOR FILING DATE: 2002-01-24
PRIOR FILING DATE: 2003-10-24
PRIOR FILING DATE: 2003-10-28
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin version 3.2
SEQ ID NO 1
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## Sequence 15, Application US/10497628

## Publication No. US20050009752A1

## SPELICANT: Teruco Nishida

## APPLICANT: Teruco Nishida

## APPLICANT: Makoto Inui

## APPLICANT: PRIOR APPLICANT: 204-06-03

## PRIOR APPLICATION NUMBER: US-001-368103

## PRIOR PILING DATE: 2001-12-01

## SOFTWARE: PatentIn Ver. 2.1

## SOFTWARE: PatentIn Ver. 2.1
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100.0%; Pred. No. 1.3e+06;
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100.0%; Pred. No. 1.3e+06;
tive 0; Mismatches 0;
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Publication No. US20040171093A1
GENERAL INFORMATION:
                                         SOFTWARE: Patentin version 3.1 SEQ ID NO 3 LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; LOCATION: (5)...(5)
; OTHER INFORMATION: AMIDATION
US-10-805-881-1
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Best Local Similarity 100.
Matches 5; Conservative
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ORGANISM: Homo sapiens
                                                                                                        ; TYPE: PRT
; ORGANISM: mammalian
US-10-688-741-3
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; ORGANISM: Human
US-10-497-628-15
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                                                         TITLE OF INVENTION: Method for Detecting Deficient Cellular Membrane Tightly Bound Ma
TITLE OF INVENTION: for Disease Diagnosis
FILE REFERENCE: N142-7005
CURRENT APPLICATION NUMBER: US/10/053,669
CURRENT FILING DATE: 2002-01-24
PRIOR FILING DATE: 1999-03-10
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin version 3.0
LENGTH: 5
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Sequence 3, Application US/10688741
Sequence 3, Application US/10688741
Sequence 3, Application US/106636A1
Sequence 3, Application OS US20040106636A1
GENERAL INFORMATION:
APPLICANT: Kream, Richard M.
APPLICANT: Kream, Richard M.
TITLE OF INVENTION: Method Of Inhibiting Opioid Tolerance Development With Chimeric 1
TITLE OF INVENTION: Analgesics
FILE REFERENCE: Kream
CURRENT APPLICATION NUMBER: US/10/688,741
CURRENT APPLICATION NUMBER: 2003-10-17
NUMBER OF SEQ ID NOS: 3
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100.0%; Pred. No. 1.3e+06;
Live 0; Mismatches 0;
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APPLICANT: Kream, Richard M.
APPLICANT: Kream, Richard M.
APPLICANT: Kream, Richard M.
TITLE OF INVENTION: Chimeric Hybrid Analgesics
FILE REFERENCE: Kream
CURRENT APPLICATION NUMBER: US/10/134,187
CURRENT FILING DATE: 2002-04-26
NUMBER OF SEQ ID NOS: 3
SOFTWARE: Patentin version 3.1
LENGTH: 5
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; Publication No. US20030202981A1
; GENERAL INFORMATION:
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; LOCATION: (5)...(5)
; OTHER INFORMATION: AMIDATION
US-10-053-669-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
Matches 5, Conservative
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GENERAL INFORMATION:
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 5; Conserv
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APPLICANT: Wells, Ibert
TITLE OF INVENTION: Method for Detecting Deficient Cellular Membrane Tightly Bound Mag
TITLE OF INVENTION: for Disease Diagnosis
FITLE OF INVENTION: for Disease Diagnosis
CURRENT APPLICATION NUMBER: US/09/265,690C
CURRENT FILING DATE: 1999-03-10
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Segure 1 Application US/10230133

Segure 3. Application US/10230133

Publication No. US20030040625A1

GENERAL INFORMATION:
APPLICANT: Wells. IDert
TITLE OF INVENTION:
TITLE OF INVENTION: methods for treatment of abnormal physiological states
TITLE OF INVENTION: methods for treatment of abnormal physiological states
TITLE OF INVENTION: methods for treatment of abnormal physiological states
CURRENT APPLICATION NUMBER: US/10/230,133

CURRENT FILING DATE: 2002-08-29

NUMBER OF SEQ ID NOS: 4

SEQ ID NO 3

LENGTH: 4

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LENGTH: 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     77.8%; Score 21; DB 9; Length 4; 100.0%; Pred. No. 1.3e+06; tive 0; Mismatches 0; Indels
                                                                                                      Sequence 2, Application US/09265690C Publication No. US20010051345A1 GENERAL INFORMATION:
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NAME/KEY: MOD RES
LOCATION: (4)...(4)
US-09-265-690C-2
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; OTHER INFORMATION: AMIDATION
US-10-230-133-3
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ORGANISM: Homo sapiens
FEATURE:
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 4; Conserva
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1 FFGL 4
                                                                  RESULT 9
US-09-265-690C-2
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US-10-230-133-3
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                                                                  Gaps
                                                                                                                                                                                                                                                                                                                        APPLICANT: Teruo Nishida
APPLICANT: Makoto Inui
APPLICANT: Makoto Inui
APPLICANT: Makoto Inui
APPLICANT: Makoto Inui
APPLICANT: Masateugu Nakamura
APPLICANT: Nasateugu Nakamura
ITILE OF INVENTION: NOVEL PEPTIDE AND PHARMACEUTICAL USE OF THE SAME
FILE REFERENCE: 04355/HG
CURRENT APPLICATION NUMBER: US/10/497,628
CURRENT FILING DATE: 2004-06-03
PRIOR APPLICATION NUMBER: JP 2001-368103
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 16
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               100.0%; Score 27; DB 17; Length 5; 100.0%; Pred. No. 1.3e+06; Ative 0; Mismatches 0; Indels
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Pred. No. 1.3e+06;
1; Mismatches 0; Indels
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APPLICANT: Benner, Robert
TITLE OF INVENTION: Gene regulator
FILE REFERENCE: 2183-5223US
CURRENT APPLICATION NUMBER: US/10/821,240A
CURRENT FILING DATE: 2004-04-08
PRIOR FILING DATE: 2001-12-21
PRIOR FILING DATE: 2001-10-04
NUMBER OF SEQ ID NOS: 312
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 270
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                                                                                                                                                                                                                                                               ; Sequence 16, Application US/10497628; Publication No. US20050009752A1; GENERAL INFORMATION:
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Best Local Similarity 80.0%;
Matches 4; Conservative
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                                                                  5; Conservative
                  Query Match
Best Local Similarity
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1 YFGLM 5
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; ORGANISM: Human
US-10-497-628-16
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US-10-497-628-16
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Sequence 2, Application US/10497628
; Publication No. US20050009752A1
; Publication No. US20050009752A1
; GENERAL INFORMATION:
; APPLICANT: Teruo Nishida
; APPLICANT: MAKOLO Inui
; APPLICANT: MAKOLO Inui
; APPLICANT: MAKOLO Inui
; TITLE OF INVENTION: NOVEL PEPTIDE AND PHARMACEUTICAL USE OF THE SAME
; FILE REPERENCE: 04155/HG
; CURRENT APPLICATION NUMBER: UP 2001-368103
; PRIOR APPLICATION NUMBER: JP 2001-368103
; PRIOR FILING DATE: 2001-12-01
; NUMBER OF SEQ ID NOS: 17
; SEO ID NO? 2
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Publication Wo. US20040142379A1
GENERAL INFORMATION:
APPLICANT: St. Hilaire, Phaedria
TITLE OF INVENTION: AFFINITY FISHING FOR LIGANDS AND PROTEIN RECEPTORS
FILE REPRENCE: 11225.16U801
CURRENT APPLICATION NUMBER: US/10/346,737A
NUMBER OF SEQ ID NOS: 50
SOFTWARE: Patentin version 3.1
SEQ ID NO 30
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Pred. No. 1.3e+06;
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                                                  TITLE OF INVENTION: Magnesium for Disease Diagnosis FILE REFERENCE: 800812-0005
CURRENT APPLICATION NUMBER: US/10/805,881
CURRENT FILING DATE: 2004-22
PRIOR APPLICATION NUMBER: US 10/053,669
PRIOR FILING DATE: 2002-01-24
PRIOR FILING DATE: 2003-01-28
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin version 3.2
SEQ ID NO 2
LENGTH: 4
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100.0%; Pred. No. ...
0; Mismatches
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; OTHER INFORMATION: AMIDATION
US-10-805-881-2
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Best Local Similarity 100.
Matches 4; Conservative
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Best Local Similarity 100.
Matches 4; Conservative
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ORGANISM: Homo sapiens
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NAME/KEY: MOD RES
LOCATION: (4) .. (4)
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ORGANISM: Human
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GENERAL INFORMATION:
APPLICANT: Wells, Ibert
TITLE OF INVENTION: Method for Detecting Deficient Cellular Membrane Tightly Bound Ma
TITLE OF INVENTION: for Disease Diagnosis
FILE REFERENCE: N1427-005
CURRENT APPLICATION NUMBER: US/10/053,669
CURRENT FILING DATE: 2002-01-24
PRIOR APPLICATION NUMBER: 09/265,690
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin version 3.0
SEQ ID NO 2
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Sequence 3. Application US/1069536

GENERAL INFORMATION:
APPLICANT: Wells, ibert Clifton
TITLE OF INVENTION: Antagonists of the Magnesium Binding Defect as Therapeutic Agents
TITLE OF INVENTION: and Methods for Treatment of Abnormal Physiological States
FILE REPERENCE: 800812-008
FILE REPERENCE: 800812-008
FILE REPERENCE: 800812-008
FILE REPERENCE: 800812-008
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FILE REPERENCE: 8008-8-29
FILE REPERENCE: 8008-89
FILE REP
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Sequence 1, Application US/10833A1
GENERAL INFORMATION:
APPLICANT: Wells, Ibert C.
APPLICANT: Wells, Method for Detecting Deficient Cellular Membrane Tightly Bound
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100.0%; Pred. No. 1.3e+06;
iive 0; Mismatches 0;
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100.0%; Pred. No. 1.38+06;
vative 0; Mismatches 0;
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NAME/KEY: MOD_RES

LOCATION: (4)...(4)

OTHER INFORMATION: AMIDATION
US-10-053-669-2
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Best Local Similarity 100.
Matches 4; Conservative
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Best Local Similarity 100.
Matches 4; Conservative
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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US-10-695-536-3
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TYPE: PRT

ORGANISM: Artificial Sequence

FEATURE:

OTHER INFORMATION: Peptide

FEATURE:

NAME/FEX: MISC_FEATURE

COTHER INFORMATION: Xaa is T(Sa)

US-10-346-737A-30

Query Match

Best Local Similarity 100.0%; Pred. No. 1.3e+06;

Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps

Oy 2 FGLM 5

Db 2 FGLM 5

Search completed: March 23, 2005, 15:07:05

JOD time: 92 secs
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Scoring table:

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Minimum DB Maximum DB

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Perfect score:

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; Sequence 2, Application US/07934553
; Patent No. 2314690
; Patent No. 2314690
; Patent No. 214690
; TITLORNATION: RATHLEEN E
; TITLE OF INVENTION: METHOD AND COMPOSITION FOR REDUCING 1GE
; TITLE OF INVENTION: MATIBODIES TO SPECIFIC ALLERGENS
; NUMBER OF SEQUENCES: 5
; CORRESPONDENCE ADDRESS:
; ADDRESSE: TILTON, FALLON, LUNGMUS & CHESTNUT
; STREET: 100 SOUTH WACKER DRIVE
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COUNTRY: USA
ZUP: G0606-4002
COMPUTER: EDADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-COMPATIBLE
OPERATING SYSTEM: PC-COMPATIBLE
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/934,553
FILING DATE: 19920821
FILING DATE: 19920821
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: US 07/705,071
FILING DATE: 24-MA-1991
ATTORNEY/AGENT INFORMATION:
NAME: FERTRESS, SUSAN B
REGISTRATION NUMBER: 31,327
REFERENCE/DOCKET NUMBER: 31,327
REFERENCE/DOCKET NUMBER: 31,327
RELEPHONE: 312/456-8000
INFORMATION FOR SEQ ID NO: 2:
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ALIGNMENTS
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Matches 5; Conservative
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    STRANDEDNESS:
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                                                                                                                                             March 23, 2005, 14:37:21; Search time 30 Seconds (without alignments) 12.442 Million cell updates/sec
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Sequence 48, P
Sequence 47, P
Sequence 49, P
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Sequence 80, A
Sequence 6, A
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Sequence 2, A
Sequence 6, A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 63
Sequence 2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 4,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Description
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PcTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PcTUS_COMB.pep:*
                        GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-07-934-553-2

US-08-225-474-2

US-09-265-690C-1

US-07-737-371E-48

US-07-737-371E-47

US-07-737-371E-47

US-08-303-362A-63

US-08-303-362A-63

US-09-635-266-3

US-09-635-266-3

US-09-635-266-3

US-09-635-266-3

US-09-635-266-3

US-08-310-3010-6

US-08-08-13-1

US-08-391-910-1

US-08-391-910-1

US-08-391-910-1

US-08-391-910-1

US-08-391-910-1

US-08-441-591-61

US-08-441-15-1

US-08-441-13-1

US-08-453-874-1
                                                                                                                                                                                                                                                                                                                                                                                                                        Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                 513545 segs, 74649064 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

    protein search, using sw model

                                                                                                                                                                                                                                                                                                                                         Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patents AA:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query
Match Length
                                                                                                                                                                                                                                                                          1 ffglm 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    seq length: 0
seq length: 5
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Gaps

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Score

Result Š.

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US-08-25-17-2

US-08-25-17-2

Sequence J. Application US/08225474

Sequence J. Application US/08225474

Sequence J. Application US/08225474

SEQUENCE J. Application US/08225474

SEQUENCE J. SETTING THE SETTING
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GENERAL INFORMATION:
APPLICANT: Wells, Ibert
TITLE OF INVENTION: Method for Detecting Deficient Cellular Membrane Tightly Bound Mac
TITLE OF INVENTION: For Disease Diagnosis
TITLE OF INVENTION: for Disease Diagnosis
FILE REFERENCE: 1427001
CURRENT APPLICATION NUMBER: US/09/265,690C
CURRENT FILING DATE: 1999-03-10
NUMBER OF SEQ ID NOS: 4
SOFTWARES PATENTIN Version 3.0
SOFTWARES PATENTIN Version 3.0
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
COUNTRY: US
ZIP: OLIO-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows5
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATE: US/07/737,371E
FILING DATE: 29-JUL-1991
CLASSIFICATION NUMBER: 07/559,172
FILING DATE: 27-JUL-1990
APPLICATION NUMBER: 07/559,172
FILING DATE: 27-JUL-1990
ATTORNEY AGENT INFORMATION:
NAME: Freeman, John W. RGGISTRANION NUMBER: 0108/028002
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: 617-542-8906
TELEFAX: 617-542-8906
TELEFX: 200154
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TUDENCAT AMINO ACIDS
TELENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ouery Match
Best Local Similarity 100.0%; Pred. No. 4.1e+05;
Matches 5; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            % Sequence 1, Application US/09265690C Patent No. 6372440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:

NAME/KEY: MOD RES

LOCATION: (5)...(5)

OTHER INFORMATION: AMIDATION
US-09-265-690C-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 100.
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear; MOLECULE TYPE: protein US-07-737-371E-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 FFGLM 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 FFGLM 5
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RESULT 5

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81.5%; Score 22; DB 2; Length 5; 100.0%; Pred. No. 4.1e+05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Yankner, Bruce A.
TITLE OF INVENTION: SCREENING METHODS TO IDENTIFY
TITLE OF INVENTION: NEUROTOXIN INHIBITORS (AS AMENDED)
NUMBER OF SQUENCES: 77
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 255 Franklin Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the LOCATION: 5...5
CTHER INFORMATION: where Xaa at position 5 is Nle US-07-737-371E-47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/737,371E
FILING DATE: 29-JUL-1991
CLASSIFCATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/559,172
FILING DATE: 27-JUL-1990
ATTOMNY AGENT INFORMATION:
NAME: Freeman, John W.
                                                                                                              CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 07/559,172
FILING DATE: 27-JUL-1990
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 29,066
REFERENCE/DOCKET NUMBER: 29,066
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: 617-542-6970
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               00108/028002
                                       US/07/737,371E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 49, Application US/0773731E Patent No. 5876948 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Freeman, John W. REGISTRATION NUMBER: 29,066
                                                                                                                                                                                                                                                                                                                                                                                                                                                           47:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 00
TELECOMMUNICATION INFORMATION
TELEPHONE: 617-542-8906
TELEFAX: 617-542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  49:
                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 617-542-890b
TELEX: 200154
INFORMATION FOR SEQ ID NO: 4'
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amind acids
      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/0
FILING DATE: 29-JUL-1991
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 100.
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Boston
STATE: MA
COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: protein FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-07-737-371E-49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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US-07-737-371E-47

Sequence 47, Application US/07737371E

Patent No. 5876948

APPLICANT: Yankner, Bruce A.
TITLE OF INVENTION:
TITLE OF INVENTION:
WUMBER OF SEQUENCES: 77

CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STRRET: 225 Franklin Street
CITY: Boston
US-07-737-371E-48
; Sequence 48 Application US/07737371E
; Patent No. 2876948
; GENERAL INFORMATION:
    TITLE OF INVENTION:
    TITLE OF INVENTION:
    NUMBER OF SEQUENCES: 77
; CORRESPONDENCE ADDRESS:
    ADDRESSEE: Fish & Richardson, P.C.
    STREET: 225 Franklin Street
    COITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    88.9%; Score 24; DB 2; Length 5; 80.0%; Pred. No. 4.1e+05; vative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: WA
COMPUTER: READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: ENGINELE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: Windows95
SOFFWARE: FASTSEE for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/737,371E
FILING DATE: 29-JUL-1991
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/559,172
FILING DATE: 27-JUL-1990
ATTORNEY/AGENT INPORMATION:
NAME: Freeman, JOHN W.
REGISTRATION NUMBER: 29,066
REFREENCE/DOCKET NUMBER: 29,066
REFREENCE/COCKET NUMBER: 29,066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM FIB: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 80.0
Elector 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  , MOLECULE TYPE: protein US-07-737-371E-48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 FFGLM 5
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Gaps

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US-08-303-362A-63
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US-08-303-362A-63
                                                                                                                                                   Query Match
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                                                                                                                   ; LOCATION: 5...5
; OTHER INFORMATION: where Xaa at position 5 is ethionine
US-07-737-371E-49
                                                                                                                                                                                                                         81.5%; Score 22; DB 2; Length 5;
100.0%; Pred. No. 4.18+05;
Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER REALABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 MG storage COMPUTER: IBM comparible OPERATING SYSTEM: MS-DOS SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/441,591
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INDORMATION:
APPLICANT: NIEUWLANDT, D., GOLD, L. AND WECKER, M.
TITLE OF INVENTION: HIGH-APPINITY
TITLE OF INVENTION: OLIGONUCLEOTIDE LIGANDS
TITLE OF INVENTION: TO THE TACHYKININ
TITLE OF INVENTION: SUBSTANCE P
NUMBER OF SEQUENCES: 66
CORRESPONDENCE ADDRESS:
ADDRESSEE: SWARBON & BRATECHUM, L.L.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Swanson & Bratschun, L.L.C.
STREET: 8400 E. Prentice Avenue, Suite 200
CITY: Englewood
STATE: Colorado
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
FILING DATE:
REGISTRICATION DATA:
REPLICATION NUMBER: 08/303,362
FILING DATE: 9-SEPTEMBER-1994
REILING DATE: 9-SEPTEMBER-1994
RELING DATE: 10-JUNE-1991
FILING DATE: 10-JUNE-1991
RELING DATE: 17-AUGUST-1992
RELING DATE: 17-AUGUST-1992
RELING DATE: 17-AUGUST-1992
RELING DATE: 17-AUGUST-1992
RELING DATE: 18-SEPTEMBER 1993
RELING DATE: 19-301
RELING DATE: 11-JUNE-1990
REILING DATE: 11-JUNE-1990
RELING DATE: 11-JUNE-1990
RELING DATE: 11-JUNE-1990
RELING DATE: 11-JUNE-1990
RELING DATE: 11-JUNE-1990
RESISTRATION NUMBER: 07/536,458
FILING DATE: 11-JUNE-1990
RESISTRATION NUMBER: 07/536,454
FILING DATE: 11-JUNE-1990
RESISTRATION NUMBER: 33,215
REFERENCE/DOCKET NUMBER: NEX21/C
TELECOMMUNICATION NUMBER: NEX21/C
                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 8
US-08-441-591-63
i Sequence 63, Application US/08441591
i Patent No. 5637682
                                                                                                                                                                                                              Query Match
Best Local Similarity 100.0
Matches 4, Conservative
5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (303) 793-3433
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS;
                             TOPOLOGY: linear MOLECULE TYPE: protein FEATURE:
                     amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 80111
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Gaps
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                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                        77.8%; Score 21; DB 1; Length 4; 100.0%; Pred. No. 4.1e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTEX: USA
ZIP: 80111
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 MG storage
COMPUTER: IBM compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/303,362A
FILING DATE: 9-SEPTEMBER-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: 07/714,131
FILING DATE: 10-JUNE-1991
FILING DATE: 10-JUNE-1991
FILING DATE: 10-JUNE-1992
FILING DATE: 17-AUGUST-1992
FILING DATE: 17-AUGUST-1992
FILING DATE: 17-AUGUST-1992
FILING DATE: 11-JUNE-1990
FILING DATE: 12-OCTOBER-1992
ATTORNEY/AGENT INFORMATION:
NAME: BATTY J. SWANSON
REGISTATION NUMBER: NEX21
FELECOMMUNICATION INFORMATION:
NAME: BATTY J. SWANSON
REGISTATION OFFER: 33-3333
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: WIEDWILANDT, D., GOLD, L. AND WECKER, M.
TITLE OF INVENTION: HIGH-AFFINITY
TITLE OF INVENTION: OLICONUCLEOTIDE LIGANDS
TITLE OF INVENTION: SUBSTANCE P
NUMBER OF SEQUENCES: 66
CORRESPONDENCES: 66
CORRESPONDENCE ADDRESS:
ADDRESSE: Swanson & Bratschun, L.L.C.
STREET: 8400 E. Prentice Avenue, Suite 200
STATE: Colorado
STATE: Colorado
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            77.8%; Score 21; DB 1; Length 4; 100.0%; Pred. No. 4.1e+05;
                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 63, Application US/08303362A; Patent No. 5648214; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4; Conservative
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-441-591-63
                                                                                                                                                                                                     Best Local Similarity
Matches 4; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
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Parent No. 664420
GENERAL INFORMATION:
APPLICANT: Wells, Ibert
TITLE OF INVENTION: Antagonists of the magnesium binding defect as therapy agents and
TITLE OF INVENTION: Antagonists of the magnesium binding defect as therapy agents and
TITLE OF INVENTION: Antagonists of treatment of abnormal physiological states
FILE REFERENCE: 2892-106
CURRENT APPLICATION NUMBER: US/10/230,133
CURRENT FILING DATE: 2002-08-29
PRIOR APPLICATION NUMBER: 09/635,266
PRIOR APPLICATION NUMBER: 09/635,266
PRIOR PILING DATE: 2000-08-09
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PatentIn version 3.0
SEQ ID NO 3
ILENGTH: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: GOLD, LARRY
APPLICANT: MISUMLANDT, DAN
APPLICANT: MISUMLANDT, DAN
APPLICANT: MISUMLANDT, DAN
APPLICANT: WECKER, MATTHEW
APPLICANT: SCHNEIDER, DANIEL J.
APPLICANT: STULENER, DANIEL J.
APPLICANT: SULLENER, BRUCE A.
APPLICANT: DOUDNA, JENNIFER, A.
TITLE OF INVENTION: HIGH-AFFINITY LIGANDS OF
TITLE OF INVENTION: P, HIV INTEGRASE AND HIV-1 REVERSE TRANSCRIPTASE
NUMBER OF SEQUENCES: 239
CORRESPONDENCE ADDRESS:
ADDRESSEE: Swanson & Bratschun, L.L.C.
STREET: 8400 E. Prentice Avenue, Suite 200
CITY: Englewood
STREET: Colorado
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MEDIUM TYPE: Diskette, 3.5 inch, 1.44 MG
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: Wordberfect 5.1
CURRENT APPLICATION DATA:
FILING DATE:
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    Sequence 3, Application US/10230133
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/23
FILING DATE: 06-MAY-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; LOCATION: (4) ... (4)
; OTHER INFORMATION: AMIDATION
US-10-230-133-3
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FILING DATE: 24-MAY-1994
CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: MOD_RES
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Patent No. 6455734

GENERAL INFORMATION:

APPLICANT: Wells, IDert

TITLE OF INVENTION: methods for treatment of abnormal physiological states

TITLE OF INVENTION: methods for treatment of abnormal physiological states

TITLE OF INVENTION: methods for treatment of abnormal physiological states

TITLE OF INVENTION: methods for treatment of abnormal physiological states

TITLE OF INVENTION: methods for treatment of abnormal physiological states

CURRENT APPLICATION NUMBER: US/09/635,266

CURRENT FILING DATE: 2000-08-09

NUMBER OF SEQ ID NOS: 4

SEQ ID NO 3

SEQ ID NO 3
                                                                                                                                                                                      US-09-265-690C-2

Sequence 2, Application US/09265690C

Sequence 2, Application US/09265690C

Sequence 2, Application US/09265690C

GENERAL INFORMATION:

APPLICANT: Wells, Ibert

TITLE OF INVENTION: for Disease Diagnosis

TITLE OF INVENTION: for Disease Diagnosis

FILE REFERENCE: 1427001

CURRENT PILING DATE: 1999-03-10

CURRENT PILING DATE: 1999-03-10

NUMBER OF SEQ ID NOS: 4

SOFTWARE: Patentin version 3.0

SEQ ID NO 2

SEQ ID NO 2
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    Mismatches
    o;
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NAME/KEY: MOD_RES
LOCATION: (4)..(4)
OTHER INFORMATION: AMIDATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CRGANISM: Homo sapiens
FEATURE:
NAME/KEY: MOD.RES
LOCATION: (4) ...(4)
OTHER INFORMATION: AMIDATION
US-09-265-690C-2
4; Conservative,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4; Conservative
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 4; Conserv
                                                   FGLM 5
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US-10-230-133-3
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, MOLECULE TYPE: peptide
US-08-070-301-6
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US-07-753-909B-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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Pred. No. 4.1e+0
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                 FILING DATE:

PRIOR APPLICATION DATA:

APPLICATION NUMBER:

PRIOR APPLICATION NUMBER:

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FILING DATE:

APPLICATION NUMBER:

TELEFROME:

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TELEFROME:

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APPLICATION NUMBER:

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APPLICATION NUMBER: 08/303,362
FILING DATE: 09-SEPTEMBER-1994
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APPLICANT: KAMINUMA, Toshihiko
APPLICANT: FUGE, Yuka
APPLICANT: TAJINA, Masahiro
APPLICANT: YANAGI, Mitsuo
APPLICANT: KRHIMOTO, Hiroshi
APPLICANT: KISHIMOTO, Jiro
APPLICANT: KISHIMOTO, Jiro
APPLICANT: KATO, Ichiro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 6, Application US/08070301
Patent No. 5871995
GENERAL INFORMATION:
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ZIP: 20036-8218
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              single
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ADDRESSEE: Wegner, C
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STRANDEDNESS: sir
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PCT-US95-05600-80
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US-08-070-301-6
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COMPUTER: INP Compatible
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COMPUTER: Percond the compatible
CLASSIFICATION NAME: Percond the compatible
FILING DARF: 13-00-189
FILING DAR
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PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:

PRILING DATE:

04-JUN-191

ATTORNEY/AGENT INFORMATION:

NAME: Sease, Edmund J.

REGISTRATION NUMBER: 24,71

FELECOMMUNICATION INFORMATION:

TELEPHONE: (515)-288-3867

TELECOMMUNICATION NUMBER: 34,71

TELECOMMUNICATION NUMBER: 36,71

TELEPAX: (515)-288-138

INFORMATION FOR SEQ ID NO. 3:

SEQUENCE CHARACTERISTICS:

LENGTH: 5 amino acids

TYPE: AMINO ACID

STRANDEDNESS: single

TOPOLOGY: linear

TOPOLOGY: linear
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Search completed: March 23, 2005, 14:50:58 Job time : 31 secs

1 FFGLM 5 | | | | 1 FXGLM 5

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RESULT 3
E4481.
2.5K structural protein - Leuconostoc
C;Species: Leuconostoc oenos phage P54
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27.5K structural p
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19 heavy chain CRD
Met-enkephalin - b
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cardioexcitatory n
hypothetical prote
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                                                                                                                                         March 23, 2005, 14:38:05; Search time 23.5 Seconds (without alignments) 20.472 Million cell updates/sec
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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Total number of hits satisfying chosen parameters:
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Maximum Match 100%
Listing first 45 summaries
                                                                                              OM protein - protein search, using sw model
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oenos phage P54 (fragment)

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C;Date: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998 C;Accession: B44817
N;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J. Gen. Microbiol. 137, 2135-2139, 1991
A;Title: Lysogeny in Leuconostoc oenos.
A;Reference number: A44817; MUID: 92085033; PMID:1748868
A;Accession: B44817
A;Molecule type: protein
A;Molecule type: protein
A;Molecule cype: protein
A;Note: sequence extracted from NCBI backbone (NCBIP:70336)
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28.5K structural protein - Leuconostoc oenos phage PAt5-12 (fragment)
28.5K structural protein - Leuconostoc oenos phage PAt5-12 (fragment)
C;Species: Leuconostoc oenos phage PAt5-12
C;Accession: C44817
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A;Ttlus: Lysogeny in Leuconostoc oenos
A;Reference number: A44817; MUID:92085033; PMID:1748868
A;Accession: C44817
A;Molacule type: protein
A;Residues: 1-5
A;Rote: sequence extracted from NCBI backbone (NCBIP:70341)
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28K structural protein - Leuconostoc oenos phage PZt11-15 (fragment)
C;Becies: Leuconostoc oenos phage PZt11-15
C;Becies: Leuconostoc oenos phage PZt11-15
C;Accesion: A44817
C;Accesion: A44817
B;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J. Gen. Microbiol. 137, 2135-2139, 1991
A;Title: Lysogeny in Leuconostoc oenos.
A;Reference number: A44817; MUID:92085033; PMID:1748868
A;Accession: A44817
A;Molecule type: protein
A;Residues: 15 <ARE>
A;Note: sequence extracted from NCBI backbone (NCBIP:70343)
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IG heavy chain CRD3 region (clone 2-100B) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
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A61445

Met-enkephalin - blue mussel

C;Species: Mytius edulis (blue mussel)

C;Species: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 21-Jan-2000

C;Accession: A6144 #sequence_revision 07-Oct-1994 #text_change 21-Jan-2000

R;Leung, M.K.; Stefano, G.B.

Proc. Natl. Acad. Sci. U.S.A. 81, 955-958, 1984

A;Title: IsoOlation and identification of enkephalins in pedal ganglia of Mytilus edulis of A;Reference number: A61445; MUID:84144823; PMID:6583690
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C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: Pr0278
B;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and jc
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44.0%; Score 11; DB 2; Length 4;
Best Local Similarity 50.0%; Pred. No. 2.8e+05;
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Best Local Similarity 20.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 3; Mismatches 1; Indels
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A53284
T-cell receptor beta 2 chain D region, Dbeta2 - rabbit
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A;Residues: 1-5 <YAM>
A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; immunoglobulin
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A,Residues: 1-5 <LEU>
A,Experimental source: pedal ganglia
C,Keywords: neuropeptide; opioid peptide
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les 2, Conservative
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cholecystokinin-5 - dog
cholecystokinin-5 - dog
NyAlternate names: CCK-5
NyAlternate names: CCK-5
Cispecies: Canis lupus familiaris (dog)
Cispecies: Canis lupus familiaris (dog)
Cispecies: A32516
Cispecies: A32516
Airitle: CCK-5: sequence analysis of a small cholecystokinin from canine brain and intest Aireference number: A32516; MUD:87153871; PMID:3826354
Airitle: CCK-5: sequence analysis of Asmall cholecystokinin from canine brain and intest Aireference number: A32516; MUD:87153871; PMID:3826354
Airitle: CCK-5: Sequence analysis of Asmall cholecystokinin from canine brain and intest Aircession: A32516
Airitle: CCK-5: Sequence analysis of Asmall cholecystokinin from canine brain and intest Aircession: A32516
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C;Comment: This peptide corresponds to the five carboxyl-terminal residues of cholecysto)
C;Superfamily: gastrin
C;Superfamily: amidated carboxyl end; neuropeptide
C;Keywords: amidated carboxyl end (Phe) #status experimental
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PT0633
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 09-Jul-2004
C;Accession: PT0633
J; Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions. A;Reference number: PT0509; MUID:91277601; PMID:1711558
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C;Date: 23-Mar-1995 #sequence_revision 05-Apr-1995 #text_change 09-Jul-2004
C;Accession 1444692
C;Accession 1444692
R;Ohta, N.; Kubota, I.; Takao, T.; Shimonishi, Y.; Yasuda-Kamatani, Y.; Minakata, H.; No Biochem. Biophys. Res. Commun. 178, 486-493, 1991
A;Title: Fulicin, a novel neuropeptide containing a D-amino acid residue isolated from the A;Reference number: A44692; MUID:91315471; PMID:1859408
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Species: Ozyatolagus cuniculus (domestic rabbit)
C;Accession: 453284
B;Harindranath, N.; Alexander, C.B.; Mage, R.G.
R;Harindranath, N.; Alexander, C.B.; Mage, R.G.
MO1. Immunol. 28, 881-888, 1991
A;Title: Evolutionarily conserved organization and sequences of germline diversity and 3,Title: Evolutionarily conserved organization
A;Accession: A53284
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C;Date: 10-Mar-1994 #sequence_revision 10-Mar-1994 #text_change 07-May-1999
C;Accession: B6:168
R;Kramer, K.J.; Felsted, R.L.; Law, J.H.
J. Blol. Chem. 248, 3021-3028, 1973
J. Blol. Chem. 248, 3021-3028, 1973
A;Title: Cocconase. V. Structural studies on an insect serine protease.
A;Reference number: A6:1168; MUID:73166540; PMID:4735570
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A;Residues: 1-5 <OHT>
A;Cross-references: UNIPROT:P35905
C;Keywords: amidated carboxyl end; D-amino acid; neuropeptide
F;2/Modified site: D-asparagine (Asn) #status experimental
F;5/Modified site: amidated carboxyl end (Val) #status experimental
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A,Residues: 1-5 <RRA>
C,Keywords: 1-14 hydrolase; serine proteinase; zymogen
F;1-5/Product: cocoonase (fragment) #status experimental <MAT>
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RESULT 15
A44955
alkanal monooxygenase (FWN-linked) (EC 1.14.14.3) alpha chain - Vibrio harveyi (fragment C.Species: Vibrio harveyi
C.Species: O3-Jun-1993 #sequence_revision 03-Jun-1993 #text_change 26-May-2000
C.Accession: A44955
R.Paquatte, O.; Tu, S.C.
Photochem. Photobiol. 50, 817-825, 1989
A;Title: Chemical modification and characterization of the alpha cysteine 106 at the Vib A;Reference number: A44955
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-5 < PAQ>
C;Keywords: FMN; luminescence; monooxygenase; oxidoreductase
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                                                                                                                                                     Query Match 32.0%; Score 8; DB 2; Length 4; Best Local Similarity 50.0%; Pred. No. 2.8e+05; Matches 1; Mismatches 0; Indels
A;Molecule type: mRNA
A;Residues: 1-4 <FEE>
A;Cross-references: UNIPROT:QBBIV7
A;Experimental source: newborn thymus, strain BALB/C
C;Keywords: T-cell receptor
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Query Match
32.0%; Score 8; DB 2; Length 5;
Best Local Similarity 50.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 1; Mismatches 0; Indels

Search completed: March 23, 2005, 14:51:54 Job time: 24.5 secs

3 GL 4 |: 2 GI 3

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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- protein search, using sw model OM protein Run on:

March 23, 2005, 14:21:29 ; Search time 112.5 Seconds (without alignments) 22.759 Million cell updates/sec

25 1 fvglm 5 Perfect score: Sequence: Title:

BLOSUM62 Scoring table:

Gapop 10.0 , Gapext 0.5

1612378 segs, 512079187 residues Searched:

. 23 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 5

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

UniProt_03:*
1: uniprot_sprot:*
2: uniprot_trembl:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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EOSI_HUMAN STANDARD; PRT; 4 AA. p02731.
p02731.
21-JUL-1986 (Rel. 01, Created)
21-JUL-2004 (Rel. 01, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Homo sapiens (Human).

SELPTE

	P19918 pseudomonas Q96at0 homo sapien P13071 citrobacter		P83073 bacillus ce P36515 saccharomyc	Q08433 rattus sp. P67857 carcinus ma	P67858 limulus pol P67859 periplaneta	P24272 vibrio fisc
UC22 MAIZE UXA4_CHLTR	DCMS_PSECH Q96AT0 BIOA_CITER	BIOB_CITFR Q99007	P83073 YLM1_YEAST	Q08433 PRCT CARMA	PRCT_LIMPO PRCT_PERAM	LUXE_VIBFI
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ALIGNMENTS

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-!- PATHWAY: Plays an important role in several metabolic pathways.
-!- SUBUNIT: Homodimer (By similarity).
-!- SIMILARITY: Belongs to the triosephosphate isomerase family.
HSC-2DPAGE; P54714; DOG.
INTERPRO; IRR000652; Triophos isomrse.
PROSITE; PS00171; TIM; PARTIAL.
Direct protein sequencing; Fatty acid biosynthesis; Gluconeogenesis; Glycolysis; Isomerase; Pentose shunt.
                                                                                                                                                                                                       TISSUE=Heart;
MEDLINE=98163340; PubMed=9504812;
Dunn M.J., Corbett J.M., Wheeler C.H.;
HSC-2DPAGE and the two-dimensional gel electrophoresis database of dog heart proteins.;
Electrophoresis 18:2795-2802(1997).
                                                                                                   Canis familiaris (Dog).
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Carnivora, Fissipedia, Canidae, Canis.
                                    01-0CT-1996 (Rel. 34, Created)
01-0CT-1996 (Rel. 34, Last sequence update)
25-0CT-2004 (Rel. 45, Last annocation update)
Triosephosphate isomerase (EC 5.3.1.1) (TIM) (Triose-phosphate
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Wed Mar 23 15:33:44 2005

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TISSUE=Skin secretion;
Wabnitz P.A., Bowie J.H., Tyler M.J., Wallace J.C.;
"Peptides from the skin glands of the Australian buzzing tree frog Litori electrica. Comparison with the skin peptides from Litoria
                                                                         Litoria rubella (Desert tree frog).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea; Hylidae;
Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TISSUE=Skin secretion, Wabnitz P.A., Bowie J.H., Tyler M.J., Wallace J.C., Powie J.H., Tyler M.J., Wallace J.C., Peptides from the skin glands of the Australian buzzing tree frog Litori electrica. Comparison with the skin peptides from Litoria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Litoria rubella (Desert tree frog).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi, Amphibia, Batrachia, Anura, Neobatrachia, Hyloidea, Hylidae, Pelodryadinae, Litoria.
                                                                                                                                                                                                                                                    Aust. J. Chem. 52:639-645(1999).

-! SUBCELLULAR LOCATION: Secreted.
-! TISSUE SPECIFICITY: Skin.
Amidation; Amphibian defense peptide; Direct protein sequencing.
MOD RES
SEQUENCE 5 AA; 630 MW; 668761F2C9A00000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Aust. J. Chem. 52:639-645(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           / Match 40.0%; Score 10; DB 1; Length 5; Local Similarity 100.0%; Pred. No. 1.6e+06; nes 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                           40.0%; Score 10; DB 1; Length 100.0%; Pred. No. 1.6e+06; Ative 0; Mismatches 0; Inde
               28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (Rel. 41, Last sequence update)
(Rel. 44, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-FEB-2003 (Rel. 41, Created)
                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 100.
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05-JUL-2004 (Re-
Rubellidin 3.2.
                                                                                                                                                                                                                                                                                                                                                                                                                     1 FV 2
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                                                           Electrin 3
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ID RE32_LITRU
AC P82073;
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P38639;
                                                                                                                                                                                                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Matches
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                                                                                                                                            Proc. Natl. Acad. Sci. U.S.A. 72:4123-4127(1975).

-1 MISCELLANBOUS: These peptides are released from mast cells in lung (and other tissues) during hypersensitivity reactions (anaphylaxis). Their activities, preferentially affecting ecsinophils, include chemotaxis, chemotactic deactivation, release of enzymes, and stimulation of the hexose monophosphate shunt. GO; GO:0006935; P:chemotaxis, IDA.
GO; GO:0006955; P:immune response, IDA.
Direct protein sequencing.
V -> A (in other partide)
                                                                                                                              of
                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=76078412; PubMed=1060093; Goetzl E.J., Austen K.F.; "Purification and synthesis of eosinophilotactic tetrapeptides of human lung tissue: identification as eosinophil chemotactic factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
NCBI_TaxID=9606;
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MEDLINE-99121193; PubMed=9461295;
Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Carcinustatin 14. Cast annotation update)
Carcinustatin 14.
Carcinus maenas (Common shore crab) (Green crab).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Plecoyemata; Brachyura;
Eubrachyura; Portunoidea; Portunidae; Carcinus.
                                                                                                                                                                                                                                                                                                                                          40.0%; Score 10; DB 1; Length 4;
100.0%; Pred. No. 1.6e+06;
Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                   V -> A (in other peptide).
/FTId=VAR_005201.
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                                                                                                                                                                                                                                                                                                            4 AA; 390 MW; 6B05B862A0000000 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
05-UUL-2004 (Rel. 44, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 AA.
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STANDARD;
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les 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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P81817;
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E103_LITRU
ID E103_LITRU
AC P82099;
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                                                         SEQUENCE
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Gaps

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Indels

Matches

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Gaps

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dioxide.
-!- CATALYTIC ACTIVITY: CO + H(2)O + A = CO(2) + AH(2).
-!- COPACTOR: Binds 1 copper(1) ion, 1 molybdenum(VI) ion and 1
molybdopterin cytosine dinucleotide (MCD) per subunit.
-!- SUBUNIT: Heterotrimer consisting of a large, a medium and a small
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wabnitz P.A., Bowie J.H., Tyler M.J., Wallace J.C.; "Peptides from the skin glands of the Australian buzzing tree frog
Litori electrica. Comparison with the skin peptides from Litoria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kraut M., Hugendieck I., Herwig S., Meyer O.;
"Homology and distribution of CO dehydrogenase structural genes in
                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea; Hylidae;
Pelodryadinae; Litoria.
NCBI_TaxID=104895;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Aust. J. Chem. 52:639-645(1999).
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUB SPECIFICITY: Skin.
Amidation; Amphibian defense peptide; Direct protein sequencing.
MOD RES 5 Histidine amide.
SEQUENCE 5 AA; 616 MW; 61F2D1A059A00000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -!- FUNCTION: Catalyzes the oxidation of carbon monoxide to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-FEB-1991 (Rel. 17, Last sequence update)
5-UUL-2004 (Rel. 44, Last annotation update)
Carbon monoxide dehydrogenase large chain (EC 1.2.99.2) (CO dehydrogenase subunit L) (CO-DH L) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pseudomonas carboxydohydrogena.
Bacteria; Proteobacteria; Alphaproteobacteria; Rhizobiales;
Bradyrhizobiaceae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   36.0%; Score 9; DB 1; Length 5; 50.0%; Pred. No. 1.6e+06; tive 1; Mismatches 0; Indel8
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Direct protein sequencing; Molybdenum; Oxidoreductase.
NON TER 4 4
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                                                                                                                                                                                            28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Blectrin 4.
                                                                                                                                            5 AA.
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                                                                                                                                            PRT;
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Arch. Microbiol. 152:335-341(1989)
                                                                                                                                                                                                                                                                                                            Litoria rubella (Desert tree frog)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=90055678; PubMed=2818128;
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Best Local Similarity 50.0
Matches 1; Conservative
                                                                                                                                            STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE=Skin secretion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=290;
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     LM 3
                                                                                                            EIO4_LITRU
ID EIO4_LITRU
AC P82100;
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P19916;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Name=cutL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     rubella.";
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     요
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                                                                                                                                                                                                                                                SEQUENCE.

TISSUB=Fibroblast;

MEDLINE=95009907; bubMed=7523108;

MEDLINE=95009907; bubMed=7523108;

MEDLINE=95009907; bubMed=7523108;

MEDLINESSONO THE TOTAL TOT
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"ILME: a waterborne pheromonal peptide released by the eggs of Sepia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bukaryota; Metazoa; Mollusca; Cephalopoda; Coleoidea; Neocoleoidea;
Decapodiformes; Sepioidea; Sepiidae; Sepia.
NCBI_TaxID=6610;
                                                                                                               Mus musculus (Mouse).
Sukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus
NCBI_TaxID=10090;
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01-OCT-1994 (Rel. 30, Created)
01-OCT-1994 (Rel. 30, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Unknown protein from 2D-PAGE of fibroblasts (P19) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=20403899; PubMed=10944467; DOI=10.1006/bbrc.2000.3286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE, SUBCELLULAR LOCATION, TISSUE SPECIFICITY, AND MASS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               40.0%; Score 10; DB 1; Length 5; 33.3%; Pred. No. 1.6e+06; Atlive 2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Direct protein sequencing; Pheromone. SEQUENCE 4 AA; 505 MW; 6B16972030000000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE 5 AA; 717 MW; 7364087043100000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            lochem. Biophys. Res. Commun. 275:217-222 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-WAR-2004 (Rel. 43, Created)
29-WAR-2004 (Rel. 43, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4 AA
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100.0%; Pred. No. 1.6
ative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pheromone peptide ILME.
Sepia officinalis (Common cuttlefish).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRT;
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Best Local Similarity 100.
Matches 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Conservative
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1 WIG 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ILME SEPOF
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28.0%; Score 7; DB 1; Length 4;

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FMRF MACNI
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MEDLINE=94286417; PubMed=7912428; DOI=10.1016/0196-9781(94)90166-X;
Madrid K.P., Price D.A., Greenberg M.J., Khan H.R., Saleuddin A.S.M.,
"FMRPamide-related peptides from the kidney of the snail, Helisoma
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MEDLINE-92195994; PubMed=1686933; DOI=10.1016/0196-9781(91)90035-N,
Evans B.D., Pohl J., Kartsonis M.A., Calabrese R.L.;
"Identification of Remide neuropeptides in the medicinal leech.";
Peptides 12:897-908(1991).
                                                                                                                                                                                                                                                                                                                    Hellsoma trivolvis (Snail).

Eukaryota, Metazoa, Annelida, Clitellata, Hirudinida, Hirudinea, Arhynchobdellida, Hirudiniformes, Hirudinidae, Hirudinidae, Hirudinidae, Hirudinidae, Hirudinidae, Hirudinidae, Hirudinidae, Hirudo.
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-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the FARP (FMRFamide related peptide)
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Eukaryota; Metazoa; Cnidaria; Anthozoa; Zoantharia; Actiniaria;
Nynantheae; Actiniidae; Anthopleura.
                                      Indels
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Pred. No. 1.6e+06;
1; Mismatches 0; Indele
. No. 1.6e+06;
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MOD RES
4 4 Phenylalanine amide.
SEQUENCE 4 AA; 582 MW; 69D40729A0000000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    group.";
Proc. Natl. Acad. Sci. U.S.A. 87:5410-5414(1990).
                                                                                                                                                                                                                            01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
                                                                                                                                                                                                     4 AA.
                                  1; Mismatches
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28-FEB-2003 (Rel. 41, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Antho-RNamide.
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               50.0%; Pred.
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         Best Local Similarity 50.0
Matches 1; Conservative
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AC PS8707;
DT 28-FEB-2003
DT 28-FEB-2003
DT 28-FEB-2003
DT 05-JUL-2004
OS Anthopleura
OC Bukaryota; M
NYMANTHESE;
OX NYMANTHESE;
OX NYMANTHESE;
OX NYMANTHESE;
NN I]
RP SEQUENCE, AN SEQUENCE, AN SEQUENCE, AN REDINESE9031
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MEDLINE=78012038; PubMed=909875;
Price D.A., Greenberg M.J.;
Purification and characterization of a cardioexcitatory neuropeptide from the central ganglia of a bivalve mollusc.";
Prep. Blochem. 7:261-281(1977).
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-!- FUNCTION: Myoactive; cardioexcitatory substance. Pharmacological activities include augmentation, induction, and regularization of
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Neuron specific.
-!- MASS SPECIFOMETRY: MW=549.3; METHOD=FAB; RANGE=1-4; NOTE=Ref.1.
PIR; A35779; A35779.
Amidation; Direct protein sequencing; Neuropeptide.
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MEDLINE=94286417; PubMed=7912428; DOI=10.1016/0196-9781(94)90166-X;
Madrid K.P., Price D.A., Greenberg M.J., Khan H.R., Saleuddin A.S.M.;
"FMRFamide-related peptides from the kidney of the snail, Helisoma
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MEDLINE=92195954; PubMed=1686933; DOI=10.1016/0196-9781(91)90035-N;
Evans B.D., Pohl J., Kartsonis M.A., Calabrese R.L.;
"Identification of RFamide neuropeptides in the medicinal leech.";
Peptides 12:897-908(1991).
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MEDLINE=90259866; PubMed=2342992; DOI=10.1016/0196-9781(90)90113-J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
05-JUL-2004 (Rel. 01, Last sequence update)
05-JUL-2004 (Rel. 01, Last sequence update)
FWRRamide (Peak C) (Cardiooxcitatory neuropeptide).
Macrocallista nimbosa (Sun-ray clam),
Narciallista nimbosa (Sun-ray clam),
Hirudo medicinalis (Medicinal leech), and
Helisoma trivolvis (Snail).
Eukaryota; Metacot, Mollusca; Bivalvia; Heteroconchia; Veneroida;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Krajniak K.G., Price D.A.; "Authentic FMRFamide is present in the polychaete Nereis virens."; Peptides 11:75-77(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE, AND SYNTHESIS.
SPECIES=M.nimbosa; TISSUE=Cerebral pedal, and Visceral ganglion;
MEDLINE=77215956; PubMed=877582;
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"Structure of a molluscan cardioexcitatory neuropeptide.";
Science 197:670-671(1977).
                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                Length 4;
                                                                                                                                                                       4 4 AA; 549 MW; 64540729A0000000 CRC64;
                                                                                                                                                  3-phenyllactic acid.
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Pred. No. 1.6e+06;
1; Mismatches 0;
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NCBI_TaxID=6594, 6353, 6421, 27815;
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50.0%;
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Best Local Similarity
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P01162;
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Int. J. Pept. Protein Res. 39:258-264(1992).

-I- FUNCTION: Neuroexcitatory peptide; increases the impulse frequency and produces a spike broadening of the identified heart excitatory and produces a spo enhances the amplitude and frequency of the heart beat. Has also an effect on several other muscles.

PIR, A32480; A33480.

D-amino acid, Direct protein sequencing; Hormone.

MOD RES

2

D-phenylalanine.

SEQÜENCE 4 AA; 408 MW; 6AADD9C81000000 CRC64;
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MEDLINE=93014529; PubMed=1399265;
IShida T., In Y., Doi M., Inoue M., Yasuda-Kamatani Y., Minakata H., Ishida T., Nomoto K.;
"Crystal Structure and molecular conformation of achatin-I (H-Gly-D-Phe-Ala-Asp-OH), an endogenous neuropeptide containing a D-amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CHARACTERIZATION.
STRAIN=Ferussac; TISSUE=Heart atrium;
STRAIN=Ferussac; TISSUE=Heart atrium;
STRAIN=Ferussac; TISSUE=Heart atrium;
Fujimoto K., Kubota I., Yasuda-Kamatani Y., Minakata H., Nomoto K.,
Yoshida M., Harada A., Muneoka Y., Kobayashi M.;
Purification of achatin-I from the atria of the African giant snail,
Achatina fulica, and ite possible function.";
Biochem. Biophys. Res. Commun. 177:847-853(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Mollusca; Gastropoda; Pulmonata; Stylommatophora;
Sigmurethra; Achatinoidea; Achatinidae; Achatina.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE, CHARACTERIZATION, AND SYNTHESIS.
STRAIN=Ferussac; TISSUE=anglion;
MEDLINE=89273551; PubMed=2597281;
MEDLINE=89273551; PubMed=2597281;
MEDLINE=89273551; PubMed=2597281;
Funase K., Sun X.P., Yongsiri A., Kim K.H., Novales-Li P.,
Novales E.T., Kanapi C.G., Takeuchi H., Nomoto K.;
"Acharin-I, an endogenous neuroexcitetory tetrapeptide from Achatina Eulica Ferussac containing a D-amino acid residue.";
Biochem. Biophys. Res. Commun. 160:1015-1020(1989):
"Growth-modulating serum tripeptide is glycyl-histidyl-lysine.", Experientia 33:324-325(1977).
--- MISCELLANEOUS: This serum tripeptide has been found to stimulate growth of some cell types and to inhibit other types in vitro.
GO: GO:0001558; P:regulation of cell growth; NAS.
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                                                                                                                                                             Direct protein sequencing.
SEQUENCE 3 AA; 340 MW; 6331E8100000000 CRC64;
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01-JUN-1994 (Rel. 29, Last sequence update)
25-OCT-2004 (Rel. 45, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 AA.
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ACH1_ACHFU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TESUE-Optic lobe;

MEDLINE-98100358; PubMed-9437704; DOI-10.1016/S0196-9781(97)00241-6;

MEDLINE-98100358; PubMed-9437704; DOI-10.1016/S0196-9781(97)00241-6;

MEDLINE-98100358; PubMed-9437704; DOI-10.1016/S0196-9781(97)00241-6;

Herry J., Favrel P., Boured-Camou B.;

Tourtleffsh, Sepia officialis.";

Peptide inhibiting the motility of the mature oviduct in the critefish, Sepia officialis.";

L. Peptides 18:1469-1474(1997).

C. I. FUNCTION: Regulatory neuropeptide with myotropic activity cargeting the distal oviduct. Inhibits the motility of the oviduct by decreasing tonus, frequency and amplitude of contractions.

C. I. MASS SPECTROMETRY: WW-229.9; METHOD-MALDI; RANGE-1-2; NOTE-Ref.1.

MOD RES

SEQUENCE 2 AA; 261 MW; 7378100000000 CRC64;
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                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Mollusca; Cephalopoda; Coleoidea; Neocoleoidea;
Decapodiformes; Sepioidea; Sepiidae; Sepia.
NCBI_TaxID=6610;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
   -!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the FARP (FMRFamide related peptide)
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                                                                                                                                                                                                                                                                                              Db .,
1.6e+06;
... 0; Indels
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                                                                                                                                                                                                                                                                                        DB 1; Length 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE, FUNCTION, MASS SPECTROMETRY, AND AMIDATION
                                                                                             PIR; A01426; ECNK.
PIR; A60418; A60418.
Amidation; Direct protein sequencing; Neuropeptide.
                                                                                                                                                                                       4 4 4 Phenylalanine amide. 4 AA; 600 MW; 69D40699A0000000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-MAR-2004 (Rel. 43, Created)
29-MAR-2004 (Rel. 43, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
25-JUL-2004 (Rel. 44, Last annotation update)
Growth-modulating peptide.
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Pred. No. 1.6e
1; Mismatches
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Schlesinger D.H., Pickart L., Thaler M.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sepia officinalis (Common cuttlefish).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRT;
                                                                                                                                                                                                                                                                                     28.0%;
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Best Local Similarity 50.0
Matches 1; Conservative
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P83570;
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SEQUENCE
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Gaps

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RESULT 14 GRWM_HUMAN

Best Loc Matches

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Search completed: March 23, 2005, 14:49:56 Job time : 112.5, secs

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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- protein search, using sw model OM protein

March 23, 2005, 14:20:03; Search time 120.5 Seconds (without alignments) 16.048 Million cell updates/sec Run on:

Title:

25 1 fvglm 5 Perfect score: Sequence:

BLOSUM62 Scoring table:

Gapop 10.0 , Gapext 0.5

2105692 seqs, 386760381 residues Searched:

45841 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 5

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

A_Geneseq_16Dec04:* 1: geneseqp1980s:* Database

geneseqp1980s:*
geneseqp1990s:*
geneseqp2000s:*
geneseqp2001s:* geneseqp2002s:* geneseqp2003as:* geneseqp2003bs:* Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

geneseqp2004s:*

SUMMARIES

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uo	Fluorinat	Fluorinat	Alpha-sub	Alpha-sub	Alpha-sub	Alpha-sub	COOH-term	Cholecyst	Cholecyst	Cholecyst	Cholecyst	Tetrapept	Substance	Neutrophi	Human tac	Mammalian	Human/rat	P antagon	C-termina	Tachykini	Octopus 1	Human bet	Substance	Tachykini	Tachykini
Description	Aab82430	Aab82431	Aar33009	Aar33008	Aar33007	Aar33010	Aaw80134	Aar54549	Aar54551	Aar54550	Aar54548	Aaw41687	Aaw99643	Aay50325	Aaw92660	Aab23028	Aab23025	Aay67576	Aab66674	Aab91428	Aab70556	Aau10880	Abb10088	Aau77847	Aau77845
ΟI	AAB82430	AAB82431	AAR33009	AAR33008	AAR33007	AAR33010	AAW80134	AAR54549	AAR54551	AAR54550	AAR54548	AAW41687	AAW99643	AAY50325	AAW92660	AAB23028	AAB23025	AAY67576	AAB66674	AAB91428	AAB70556	AAU10880	ABB10088	AAU77847	AAU77845
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Score	25	25	20	20	20	20	20	20	20	20	20	20	20	20	20	20	20	20	20	20	20	20	20	20	20
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Adc64000 Tachykini Ade94203 High acti Adf92530 Substance Adm12180 Tachykini	Mammal Mammal Human Tetrap SEC bi		Aabb1821 Amyloid b Aabb1794 Amyloid b Aap20392 Analgesic Aaw92702 Human tac Aag99346 Atypical
ADC64000 ADE94203 ADF92530 ADN12180	ADN95078 ADR43774 ADR43771 AAR41684	AAB26299 AAG80461 AAW41685 AAB91795 AAB91822	AAB91821 AAB91794 AAP20392 AAW92702 AAG99346
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ALIGNMENTS

AAB82430 standard; peptide; 5 AA. RESULT 1

AAB82430;

22-AUG-2001 (first entry)

Fluorinated neurokinin A antagonist intermediate.

Neurokinin A, fluorinated peptide; antagonist; immunosuppressive; antiarthritic; antiasthmatic; antiinflammatory; antiarthritic; analgesic; antitumour; anticonvulsant; nootropic; antipsychotic; antimigraine; asthma; therapy.

Synthetic.

Location/Qualifiers 1. .2 /note= "the peptide bond is replaced by CH2-N(CH2CF3)" /note= "C-terminal amide" 'note= "Boc-Phe" 96US-00638407 Key Modified-site Modified-site Modified-site US6218364-B1 17-APR-2001. 26-APR-1996;

88US-00208926. 89US-00315202. 89US-00356031. 91US-000686593. 93US-00033987. 94US-00282341. 23-MAY-1989; 17-APR-1991; 31-MAY-1991; 19-MAR-1993; 29-JUL-1994; 20-JUN-1988; 24-FEB-1989;

(HARB/) HARBESON S L. (MCCA/) MCCARTHY J R.

я, ; Harbeson SL, Mccarthy

WPI; 2001-366135/38.

New peptide derivative useful as immunosuppressants and in treating subjects with various conditions, e.g. asthma.

Wed Mar 23

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Example 3; Col 20; 25pp; English.
                                                                 Local Similarity 100.
                                                                           1 FVGLM 5
                                                                                  1 FVGLM 5
                                                                                                                                        asthma; therapy
                                                       Sequence 5 AA;
                                                                                                                                                      Key
Modified-site
                                                                                                                                                                Modified-site
                                                                                                                 22-AUG-2001
                                                                                                                                                                           US6218364-B1
                                                                                                                                                                                        26-APR-1996;
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31-MAY-1991;
                                                                                                                                                                                                      23-MAY-1989
                                                                                                                                                                                                                19-MAR-1993
                                                                                                          AAB82431;
                                                                                                                                                Synthetic.
                                                             Query Match
                                                                Best Loc
Matches
                                                                                            RESULT 2
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The present sequence is that of a synthetic peptide produced as an intermediate in the chemical synthesis of a novel fluorinated neurokinin h antagonist (AAB82428). Fluorinated neurokinin A antagonists of the invention are based on the amino acid sequence of neurokinin A, but include at least 1 modified peptide bond having a reduced amide and a fluorinated alkyl attached to the N atom of the modified peptide bond. Treating subjects, including humans, with various conditions, e.g. asthma (claimed), arthritis, urinary incontinence, pain, inflammation, tumour growth, gastrointestinal hypermotility, Huntington's disease, psychosis, neuralgia, urticaria, carcinoid syndrome symptoms, influenza, common cold, and headache including migraine
                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 25; DB 4; Length 5; 100.0%; Pred. No. 1.8e+06; tive 0; Mismatches 0; Indels
                      Example 3; Col 20; 25pp; English.
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Best Local Similarity
Matches 5; Conserv
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                                      The present sequence is that of a protected peptide produced as an intermediate in the chemical synthesis of a novel fluorinated neurokinin h antagonist (AABS2428) Fluorinated neurokinin A antagonists of the invention are based on the amino acid sequence of neurokinin A, but include at least 1 modified peptide bond having a reduced amide and fluorinated alkyl attached to the N atom of the modified peptide bond. Treating subjects, including humans, with various conditions, e.g. asthma (claimed), arthritis, urinary incontinence, pain, inflammation, tumour prowth, gastrointestinal hypermotility, Huntington's disease, psychosis, neuralgia, urticaria, carcinoid syndrome symptoms, influenza, common cold, and headache including migraine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Neurokinin A, fluorinated peptide, antagonist, immunosuppressive; antiarthritic; antiasthmatic; antiinflammatory; antiarthritic; analgesic; antitumour; anticonvulsant; nootropic; antipsychotic; antimigraine;

    .2
/note= "the peptide bond is replaced by CH2-N(CH2CF3)"

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100.0%; Pred. No. 1.8e+06;
ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB82431 standard; peptide; 5 AA.
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91US-00709092.
93US-00033987.
94US-00282341.
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89US-00356031,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HARB/) HARBESON S L. (MCCA/) MCCARTHY J R.
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Gaps

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The peptide is a specifically claimed example of a group of generically claimed mono-, di-, tri-, tetra- and penta-peptides which include a substituent on an alpha-C atom in the chain. Such substitution may modify
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                                                                                                                                                                                                                                                                                                                                                                           CCK; neuropeptide; endorphin; hormone; LHRH; contraception; analgesia;
improved bioavailability.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Howson W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Richardson RS,
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AAR33009 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                        Alpha-substituted short peptide.
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New peptide derivative useful as immunosuppressants and in treating subjects with various conditions, e.g. asthma.

Harbeson SL, Mccarthy JR;

29-JUL-1994;

WPI; 2001-366135/38.

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the bioavailability, stability or absorbability of the peptide and hence may improve the activity of the peptide as a drug. Depending on the nature of the parent peptide (hormone, endorphin, CCK, NK2, chemotactic peptide, etc.), the modified peptides are variously useful for treating obesity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, addictive drug withdrawal symptoms, hypertension, heart failure, cognition or memory disorders, spasitity, depression, diabetes, cancer, asthma, bladder dysfunction, psychosis and arthritis; and as contracoptives. (Updated on 25-MAR-2003 to correct PD field.) (Updated on 25-MAR-2003 to correct PR field.)
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Sequence 5 AA;

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Length 5;
                       1; Indels
Score 20; DB 2; I
Pred. No. 1.8e+06;
0; Mismatches 1;
 80.0%;
80.0%;
                       4; Conservative
Query Match
Best Local Similarity
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                        Matches
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Gaps ö

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AAR33008 standard; peptide; 5 AA.
            (first entry)
          (revised)
          25-MAR-2003
02-APR-1993
       AAR33008;
 RESULT 4
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Alpha-substituted short peptide

CCK; neuropeptide; endorphin; hormone; LHRH; contraception; analgesia; improved bioavailability.

Synthetic

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'note= "alpha-Me-Phe"
Location/Qualifiers
                                                                 /note= "Met-NH2
               Modified-site
                                               Modified-site
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W09219254-A1

12-NOV-1992

92WO-US003119. 92US-00852086 91US-00690755 15-APR-1992; 24-APR-1991; 20-MAR-1992;

(WARN) WARNER LAMBERT CO

Howson W; Richardson RS, Horwell DC, Hughes J,

WPI; 1992-398522/48.

New alpha-substd. polypeptide are e.g. selective receptor ligands - for treating inflammation, pain, stroke, ulcers, hypertension, heart failure, depression, cancer, asthma, psychosis, arthritis, etc.

Claim 3; Page 41; 46pp; English.

The peptide is a specifically claimed example of a group of generically claimed mono-, di-, tri-, tetra- and penta-peptides which include a substituent on an alpha-C atom in the chain. Such substitution may modify the bicavailability, stability or absorbability of the peptide and hence may improve the activity of the peptide as a drug. Depending on the nature of the parent peptide (hormone, endorphin, CCK, NK2, chemotactic peptide, etc.), the modified peptides are variously useful for treating

The peptide is a specifically claimed example of a group of generically claimed mono-, di-, tri-, tetra- and penta-peptides which include a substituent on an alpha-C atom in the chain. Such substitution may modify the bioavailability, stability or absorbability of the peptide and hence may improve the activity of the peptide as a drug. Depending on the nature of the pract peptide (hormone, endorphin, CCK, NK2, chemotactic peptide, etc.), the modified peptides are variously useful for treating obesity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, addictive drug withdrawal symptoms, hypertension, heart failure, cognition or memory disorders, spasticity, depression, diabbetes, cancer, asthma, bladder dysfunction, psychosis and arthritis; and as

New alpha-substd. polypeptide are e.g. selective receptor ligands - for treating inflammation, pain, stroke, ulcers, hypertension, heart failure, depression, cancer, asthma, psychosis, arthritis, etc.

Claim 3; Page 41; 46pp; English

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obesity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, addictive drug withdrawal symptoms, hypertension, heart failure, cognition or memory disorders, spasticity, depression, diabetes, cancer, asthma, bladder dysfunction, psychhosis and arthritis; and as contraceptives. (Updated on 25-MAR-2003 to correct PM field.) (Updated on 25-MAR-2003 to correct PR field.) (indated on 25-MAR-2003 to correct PR field.)
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                                                                                                     Score 20; DB 2; Length 5;
Pred. No. 1.8e+06;
0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                    /note= "alpha-Me-Phe"
                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                    AAR33007 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                          Alpha-substituted short peptide.
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92US-00852086.
                                                                                                      80.0%;
80.0%;
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                                                                                                                          4; Conservative
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                                                                                                      Query Match
Best Local Similarity
Matches 4; Conserv
                                                                                                                                             1 FVGLM 5
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                                                                                    Sequence 5 AA;
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20-MAR-1992;
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02-APR-1993
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                                                                                                                                                                                                                                                                                                                                             Synthetic
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contraceptives. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PD field.) (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PR
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improved bioavailability.
                                                                      Gaps
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                                               Query Match 80.0%; Score 20; DB 2; Length 5; Best Local Similarity 80.0%; Pred. No. 1.8e+06; Matches 4; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                     /note= "alpha-Me-Met-NH2"
                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                     AAR33010 standard; peptide; 5 AA.
                                                                                                                                                                                            Alpha-substituted short peptide.
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                                                                               1 FVGLM 5
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                                 Sequence 5 AA;
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Modified-site
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02-APR-1993
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New alpha-substd. polypeptide are e.g. selective receptor ligands - for treating inflammation, pain, stroke, ulcers, hypertension, heart failure, depression, cancer, asthma, psychosis, arthritis, etc. Claim 3; Page 41; 46pp; English.

Horwell DC, Hughes J, Richardson RS,

WPI; 1992-398522/48

The peptide is a specifically claimed example of a group of generically claimed mono. 'd-' tri-, tetra- and penta-peptides which include a substituent on an alpha-C atom in the chain. Such substitution may modify the bioavailability, stability or absorbability of the peptide and hence mature of the parent peptide (hormone, endorphin, ccx, NRZ, chemotractic peptide, etc.), the modified peptides are variously useful for treating obstity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, addictive drug withdrawal symptoms, hypertension, heart failure, cognition or memory disorders, spassicity, depression, diabetes, cancer, contraceptives. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PR field.) (Updated on 25-MAR-2003 to correct PR field.)

Sequence 5 AA;

1 FXGLM

RESULT 8 AAR54549

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents the COOH-terminal sequence of the tachykinin family. The specification describes a human recombinant neurokinin (NK-2) receptor protein. The human NK-2 receptor gene was closed from human tracheal tissue from an individual with cystic fibrosis. The coding sequence is interrupted by four introns. The protein between neurokinin A neurotransmitter and its NK-2 receptor. The protein is thus useful for treating disorders associated with abnormal smooth muscle cell contraction, particularly asthma and gastrointestinal disorders such as peptic ulcers and ulcerative colitis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human recombinant neurokinin NK-2 receptor - antagonises interaction of neurokinin A and its receptor, useful for treating asthma and ulcerative
                                     Gaps
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                                                                                                                                                                                                                                                        Human, neurokinin receptor; NK-2; neurokinin A neurotransmitter; abnormal smooth muscle cell contraction; asthma; PCR primer; gastrointestinal disorder; peptic ulcer; ulcerative colitis.
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0
     Score 20; DB 2; Length 5;
Pred. No. 1.8e+06;
0; Mismatches 1; Indels
                                1; Indels
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80.0%; Pred. No. 1.8e+06;
live 0; Mismatches 1; Indels
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                                                                                                                                                                                                                              COOH-terminal sequence of the tachykinin family.
                                                                                                                                                                                                                                                                                                                                                                   /note= "Phe, Tyr, Val or Ile"
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                                                                                                                                               AAW80134 standard; peptide; 5 AA.
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    80.0%;
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                              4; Conservative
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Query Match
Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
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                                                                                                                                                                                                     17-DEC-1998
                                                                                                                                                                                                                                                                                                             Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-MAR-1991;
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AAR54549;

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The sequences given in AAR53117-38 and AAR54530-51 are peptide analogues of cholecystokinin (CCK) which can be used to treat obesity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, hypertension, heart failure, cognition, memory enhancement, spasticity, depression, diabetes, cancers, asthma, bladder dysfunction, psychosis, arthritis and in the treatment of substance withdrawl. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptide analogue; peptoid; cholecystokinin; CCK; obesity; anxiety; gastrointestinal ulcers; pain; stroke; inflammation; hypertonsion; heart failure; cognition; memory enhancement; spasticity; depression; diabetes; cancers; asthma; bladder dystunction; psychosis; arthritis.
gastrointestinal ulcers; pain; stroke; inflammation; hypertension;
heart failure; cognition; memory enhancement; spasticity; depression;
diabetes; cancers; asthma; bladder dysfunction; psychosis; arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                New cholecystokinin analogues – useful e.g. in treatment of pain, obesity, stroke, anxiety, and gastrointestinal ulcers.
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80.0%; Pred. No. 1.8e+06;
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                                                                                                          Location/Qualifiers
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                                                                                                                                           /label= MeMet
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14-DEC-1994
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                                                                                                                                                        Peptide analogue; peptoid; cholecystokinin; CCK; obesity; anxiety; gastrointestinal ulcers; pain; stroke; inflammation; hypertension; heart failure; cognition; memory enhancement; spasticity; depression; diabetes; cancers; asthma; bladder dysfunction; psychosis; arthritis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New cholecystokinin analogues - useful e.g. in treatment of pain, obesity, stroke, anxiety, and gastrointestinal ulcers.
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Pred. No. 1.8e+06;
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                                                                                                                         Cholecystokinin analogue peptide #42.
                                                                                                                                                                                                                                                                                     Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR54551 standard; peptide; 5 AA.
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 AAR54549 standard; peptide; 5
                                                                                                                                                                                                                                                                                                                    /label= MePhe
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80.0%;
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93US-00131693
                                                                     (revised)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (revised)
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                                                                                                                                                                                                                                                                                     Key
Modified-site
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08-OCT-1993;
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                                                                     25-MAR-2003
14-DEC-1994
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AAR54551;

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RESULT 9 AAR5455

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Wed Mar 23

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Claim 3; Page 66; 73pp; English
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Best Local Similarity 80.vs,
 (WARN ) WARNER LAMBERT CO.
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                                                                WPI; 1994-151243/18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   dry eye, keratitis.
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Modified-site
                                Horwell DC,
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                                                                                                                                                                                                                                                                                                         The sequences given in AARS3117-38 and AARS4530-51 are peptide analogues of cholecystokinin (CCK) which can be used to treat obesity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, hypertension, heart failure, cognition, memory enhancement, spasticity, depression, diabetes, cancers, asthma, bladder dysfunction, psychosis, arthritis and in the treatment of substance withdrawl. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Peptide analogue; peptoid; cholecystokinin; CCK; obesity; anxiety; gastrointestinal ulcers; pain; stroke; inflammation; hypertension; heart failure; cognition; memory enhancement; spasticity; depression; diabetes; cancers; asthma; bladder dysfunction; psychosis; arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                 New cholecystokinin analogues - useful e.g. in treatment of pain, obesity, stroke, anxiety, and gastrointestinal ulcers.
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80.0%; Pred. No. 1.8e+06;
iive 0; Mismatches 1; Indels
                                                                                                                                                                      Howson W, Hugues J, Richardson RS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "Amidated C-terminal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cholecystokinin analogue peptide #41.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR54548 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                             Claim 3; Page 66; 73pp; English.
                                                                93WO-US009809
                                                                                            92US-00963169.
93US-00131693.
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Best Local Similarity
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WO9409031-A1
                                                           14-OCT-1993;
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Modified-site
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                             28-APR-1994
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14-DEC-1994
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08-OCT-1993;
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Matches

RESULT 11 AAR54548

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The sequences given in AAR53117-38 and AAR54530-51 are peptide analogues of cholecystokinin (CCK) which can be used to treat obseity, anxiety, gastrointestinal ulcers, pain, stroke, inflammation, hypertension, heart failure, cognition, memory enhancement, spasticity, depression, diabetee, cancers, asthma, bladder dysfunction, psychosis, arthmis and in the treatment of substance withdrawl. (Updated on 25-MAR-2003 to correct PN field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ophthalmic drug; corneal disorder; ulcer; epithelial peeling; dry eye; keratitis; insulin like growth factor-1; IGF-1; eye drop.
                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ophthalmic drug composition containing tetra:peptide - is useful as corneal disorder remedy for corneal ulcer, corneal epithelial peeling,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This sequence is shown in the specification. The invention relates to ophthalmic drug composition which contains Phe-Gly-Leu-Met-NH2 or its medicinally acceptable salts as the active ingredient. It is used, together with insulin like growth factor-I (IGF-I), to treat corneal
                                                                    New cholecystokinin analogues - useful e.g. in treatment of pain, obesity, stroke, anxiety, and gastrointestinal ulcers.
                                                                                                                                                                                                                                                                                                                                                                                                ·.
                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                                                           Length 5;
   Richardson RS;
                                                                                                                                                                                                                                                                                                                                                    80.0%; Score 20; DB 2; I
80.0%; Pred. No. 1.8e+06;
                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
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Howson W, Hugues J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nishida T, Nakamura M, Nakata K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW41687 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CCCCCXXXIXXBXXBXXBXXBXXBXXBXXBXXBXCCCCCCCXXXIXXBXXBXXBXXBXXBX
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Gaps

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A composition has been developed for supplying a peptide to an opioid receptor or that interferes with binding of substance P to its receptor. The composition comprises: (a) myogenic cells that contain heterologous DNA encoding the peptide to express the peptide; and (b) a pharmaceutically acceptable carrier. The composition is useful for relieving pain and for treating behavioural and perceptive abnormalities using myoblast transfer therapy. It is useful in a method for treating psychiatric conditions that involve abnormal perception e.g. depression, chronic maxiety syndromes, paranoia, alcoholism and drug addiction, chronic pain and other diseases in which opioid neurons and substance P sensitive neurons play a role. The composition provides a continuous, long term supply of opioid peptides (Long-term analgesia) which lasts for up to at least 6 years. The present sequence represents a specifically
keratitis. The dosage is 0.1-5000 (preferably 1-1000) mg/day of the active ingredient and 0.001-100 (preferably 0.01-10) mg/day of the preferable form of the composition is eye drops
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New composition for supplying peptide to opioid receptor - comprises myogenic cells containing heterologous DNA encoding peptide and carrier.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Substance P; myoblast transfer therapy; pain relief; analgesic; behavioural abnormality; perceptive abnormality; opioid receptor; psychiatric condition; depression; chronic anxiety syndrome; paranoia; alcoholism; drug addiction; chronic pain; neuron.
                                                                                                                          Score 20; DB 2; Length 5;
Pred. No. 1.8e+06;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                 AAW99643 standard; peptide; 5 AA
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                                                                                                                       80.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                     Substance P analogue peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98EP-00201068
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                                                                                                                                                            Conservative
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Best Local Similarity
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Synthetic.
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80.0%; Score 20; DB 2; I 80.0%; Pred. No. 1.8e+06; iive 0; Mismatches 1;

Conservative

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cell activating compositions which involves preparing a cell activating composition compositions which involves preparing a cell activating composition compositions which involves preparing a cell activating composition comprising (a) homogenizing panceasic; (b) removing composition comprising (a) homogenizing panceasic; (b) removing the resulting particulates from the homogenate; (c) optionally incubating the resulting factionating the homogenate and selecting fractions that exhibit cell activation activity. The methods can be used for improving treatment controme or reducing risk of treatment of e.g. cardiovascular disease, inflammatory disease, trauma, autoimmune diseases, arthritis, organ rejection, diabetes and diabetic complications, stroke, ischemia, critical adheres and diabetic complications, stroke, ischemia.

They can be used in the veterinary treatment of a non-human subject. They can be used to lower cell activation resulting from the used in the veterinary treatment of a non-human subject. Protease inhibitors can be used to lower cell activation resulting from these diseases and deficiencies. The detection of an elevated level of hydrogen peroxide in plasma or whole blood and in the elevated level of hydrogen peroxide in plasma or whole blood and in the cell indicative of the onset of an acute cardiovascular disorders, such as disease onset or ischemic complications. An elevated level of hydrogen complications. An elevated level of hydrogen complication or sepsies. Any septement peptides used in the hyperension or sepsies. Appresence of substruction and in the presence of substruction or immune compromised condition e.g. method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and treatment of e.g. cardiovascular, inflammatory, autoimmune or
Alzheimer's disease, trauma, arthritis, organ rejection, diabetes, stroke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of cell activating compositions in developing products for diagnosis
                                                                                                                                                                                                                                                                                                                            Cell activation; pancreas; treatment; cardiovascular disease; trauma; inflammatory disease; autoimmune disease; arthritis; diabetes; stroke; organ rejection; ischemia; Alzheimer's disease; myocardial infarction; haemorrhagic shock; diabetic retinopathy; venous insufficiency; angina; trauma; protease inhibitor; hypertension; sepsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention describes a novel method for the use and preparation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kistler E;
                                                                                                                                                                                                                                                                                      Neutrophil-activating pancreatic derived peptide 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hugli TE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stoughton RB, Schmid-Schonbein GW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 9; Page 184; 184pp; English
                                                                                                                                                      AAY50325 standard; peptide; 5 AA
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                                                                                                                                                                                                                                          (first entry)
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1 FVGLM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unidentified
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Gaps

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Tachykinin agonist; beta-amyloid; inhibition; neurotoxin; treatment; Alzheimer's disease; Down's syndrome; amyloidosis; human; hereditary cerebral haemorrhage; non-inherited congophilic angiopathy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Screening for neurotoxin inhibitors - by testing compounds for their effect on beta-amyloid peptide neurotoxic effect on neuronal cells.
                                                                                                                                                                                                                                                  Human tachykinin agonist beta-amyloid peptide fragment #6.
                                                                          80.0%; Score 20; DB 2; Length 5;
80.0%; Pred. No. 1.8e+06;
iive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Col 13-14; 28pp; English.
                                                                                                                                                                                AAW92660 standard; peptide; 5 AA.
Wed Mar 23 15:33:42 2005
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                                                                                                                                                                                                                                                                                                                                                                                         90US-00559173.
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(first entry)
                                                                     Query Match
Best Local Similarity 80.0
Matches 4; Conservative
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                                                                                                             1 FVGLM 5
                                                                                                                                 1 FFGLM 5
                                                     Sequence 5 AA;
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                                                                                                                                                                                                                      20-MAR-2003
30-APR-1999
                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                       27-JUL-1990;
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AAW92660
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This invention describes a method for screening compounds for inhibiting a neurotoxin. The method involves incubating tachykinin agonists with neuronal calls and a betra-amyloid peptide neurotoxin. The methods can be used for identifying compounds for treating diseases characterised by an Undesirable build up of beta-amyloid protein, e.g. Alzheimer's disease, bown's syndrome, and the syndromes of hereditary cerebral haemorrhage with amyloidosis and non-inherited congophilic angiopathy with cerebral beta-amyloid peptide fragments. (Updated on 20-MAR-2003 to correct printed).

Sequence 5 AA;

0; Gaps Score 20; DB 2; Length 5; Pred. No. 1.8e+06; 0; Mismatches 1; Indels Query Match 80.0%; Best Local Similarity 80.0%; Matches 4; Conservative (

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Search completed: March 23, 2005, 14:46:01 Job time : 121.5 secs

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GenCore version 5.1.6
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OM protein - protein search, using sw model

March 23, 2005, 14:50:07; Search time 92 Seconds (without alignments) 17.995 Million cell updates/sec

Run on:

SEQ2 25 1 fvglm 5 Title: Perfect score: Sequence: **BLOSUM62** Scoring table:

Gapop 10.0 , Gapext 0.5

Total number of hits satisfying chosen parameters: 1407402 segs, 331100923 residues Searched:

Minimum DB seq length: 0 Maximum DB seq length: 5

Post-processing: Minimum Match 0% Maximum Match 100%

Listing first 45 summaries

Database :

Published Applications AA:*

1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US07_REW PUB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US08_REW PUB.pep:*

7: /cgn2_6/ptodata/1/pubpaa/US08_REW PUB.pep:*

8: /cgn2_6/ptodata/1/pubpaa/US08_REW PUB.pep:*

9: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

10: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

11: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

11: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

13: /cgn2_6/ptodata/1/pubpaa/US08_REW PUB.pep:*

14: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*

15: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*

16: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*

17: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*

18: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*

19: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*

10: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Query Match Length DB	DB	ID	Description
	20	80.0		9	US-09-265-690C-1	Sequence 1, Appli
7	20	80.0	ī.	σ	US-09-265-690C-4	Sequence 4, Appli
٣	20	80.0	S	14	US-10-230-133-4	Sequence 4, Appli
4	20	80.0	S	14	US-10-053-669-1	Sequence 1, Appli
2	20	80.0	S	14	US-10-053-669-4	Sequence 4, Appli
9	20	80.0	Ŋ	15	US-10-134-187-3	Sequence 3, Appli
7	20	80.0	S	16	US-10-688-741-3	'n
80	20	80.0	S	19	US-10-695-536-4	Sequence 4, Appli
6	20	80.0	Ŋ	76	US-10-805-881-1	H
10	20	80.0	S	16	US-10-805-881-4	4
11	20	80.0	Z,	17	US-10-497-628-15	Sequence 15, Appl
13	17	68.0	Ŋ	14	US-10-168-789A-32	Sequence 32, Appl
13	17	68.0	S	17	US-10-497-628-16	16

Sequence 30, Appl Sequence 2, Appli Sequence 14, Appl Sequence 39, Appl Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 3, Appli Sequence 3, Appli Sequence 2, Appli Sequence 270, Appli Sequence 270, Appli Sequence 3, Appli Sequence 3, Appli	30, 17, 113, 10, 10, 10, 10, 10, 10, 10, 10, 10, 10
US-08- US-10 US-10 US-10 US-10 US-10 US-10 US-10 US-10 US-10 US-10 US-10 US-10 US-10	16 US.10-346-737A-30 17 US-10-497-628-17 17 US-10-641-286-13 9 US-09-943-123-24 14 US-10-087-402-10 14 US-10-361-290-12 17 US-10-712-359A-24 17 US-10-712-359A-24 13 US-10-014-716-28 14 US-10-190-951-28 14 US-10-190-951-28 15 US-10-1599-665-15 15 US-10-436-549-565 16 US-10-436-549-565
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ALIGNMENTS

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Subjection US/09265690C
Publication US/09265690C
Publication No. US20010051345A1
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: for Disease Diagnosis
TITLE OF INVENTION: for Disease Diagnosis
TITLE OF INVENTION: for Disease Diagnosis
CURRENT APPLICATION NUMBER: US/09/265,690C
CURRENT FILING DATE: 1999-03-10
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 20; DB 9; Length 5;
Pred. No. 1.3e+06;
0; Mismatches 1; Indels
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Best Local Similarity 80.0%;
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: MOD RES;
LOCATION: (5)...(5)
CTHER INFORMATION: AMIDATION
US-09-265-690C-1
                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
US-09-265-690C-1
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1 FVGLM 5 1 FFGLM 5 RESULT 2 US-09-265-690C-4 ઠે g

; Sequence 4, Application US/09265690C; Publication No. US20010051345A1

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Gaps

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APPLICANT: Wells, lbert
TITLE OF INVENTION: for Disease Diagnosis
FITLE OF INVENTION: for Disease Diagnosis
FILE REFERENCE: N1427-005
CURRENT APPLICATION NUMBER: US/10/053,669
CURRENT FILING DATE: 2002-01-24
PRIOR APPLICATION NUMBER: 09/265,690
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PATENT VERSION 3.0
SEQ ID NO 1
LENGTH: 5
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TITLE OF INVENTION: Method for Detecting Deficient Cellular Membrane Tightly Bound Mac
TITLE OF INVENTION: for Disease Diagnosis
FILE REFERENCE: N1427-005
CURRENT APPLICATION NUMBER: US/10/053,669
CURRENT FILING DATE: 2002-01-24
PRIOR PILING DATE: 1999-03-10
NUMBER OF SEQ ID NOS: 4
SEQ ID NO 4
LENGTH: 5
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80.0%; Score 20; DB 14; Length 5;
Best Local Similarity 80.0%; Pred. No. 1.3e+06;
Matches 4; Conservative 0; Mismatches 1; Indels
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COCATION: (5)..(5)

O'THER INFORMATION: AMIDATION

NAME/KEY: VARIANT

LOCATION: (2)..(2)

O'CHER INFORMATION: "X" may be either Phe or Val.

US-10-053-669-4
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Publication No. US20030077658A1
GENERAL INFORMATION:
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Publication No. US20030202981A1
GENERAL INFORMATION:
APPLICANT: Kream, Richard M.
                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: MOD_RES

LOCATION: (5)...(5)

OTHER INFORMATION: AMIDATION

US-10-053-669-1
                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 4; Conserv
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Sequence 4, Application US/10230133
Sequence 4. Application US/10230133
Publication No. US20030040625A1
GENERAL INPORMATION:
TITLE OF INVENTION: Antagonists of the magnesium binding defect as therapy agents and TITLE OF INVENTION: methods for treatment of abnormal physiological states
CURRENT FILING DATE: 2002-08-29
PRIOR APPLICATION NUMBER: US/10/230,133
CURRENT FILING DATE: 2000-08-09
PRIOR FILING DATE: 2000-08-09
NUMBER OF SEQ ID NOS: 4
SOUTHER PRESENT OF SEQ ID NOS: 4
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80.0%; Pred. No. 1.3e+06;
ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Indels
                                                                                                                                                                                                                                                                                                TYPE: PRT

GRGANISM: Homo sapiens

GRGANISM: Homo sapiens

FRATURE:

NAME/KEY: MOD_RES

LOCATION: (5)...(5)

OTHER INFORMATION: AMIDATION

NAME/KEY: VARIANT

LOCATION: (2)...(2)

OTHER INFORMATION: "X" may be either Phe or Val.

US-09-265-690C-4
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US-10-053-669-1
; Sequence 1, Application US/10053669
; Publication No. US20030077658A1
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LOCATION: (5) (5)
OTHER INFORMATION: AMIDATION
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Best Local Similarity 80.0
Matches 4; Conservative
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Best Local Similarity 80.0
Matches 4; Conservative
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ORGANISM: Homo sapiens
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LENGTH: 5
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Sequence 4, Application US/10805881
Publication No. US20040171093A1
GENERAL INFORMATION:
APPLICANT: Wells, Ibert C.
TITLE OF INVENTION: Method for Detecting Deficient Cellular Membrane Tightly Bound
FILE OF INVENTION: Magnesium for Disease Diagnosis
FILE REPERBNCE: 800812-0005
CURRENT APPLICATION NUMBER: US/10/805,881
CURRENT APPLICATION NUMBER: US 10/053,669
PRIOR APPLICATION NUMBER: US 10/053,669
PRIOR APPLICATION NUMBER: US 10/053,669
PRIOR PLING DATE: 2002-01-24
PRIOR PLING DATE: 2003-01-24
PRIOR PLING DATE: 2003-10-28
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Wells, Ibert C.

TITLE OF INVENTION: Method for Detecting Deficient Cellular Membrane Tightly Bound
TITLE OF INVENTION: Magnesium for Disease Diagnosis
FILE OF INVENTION: Magnesium for Disease Diagnosis
FILE REPERBNCE: 800812-0005
CURRENT APPLICATION NUMBER: US/10/805,881
CURRENT APPLICATION NUMBER: US 10/053,669
PRIOR APPLICATION NUMBER: US 10/695,536
PRIOR FILING DATE: 2003-10-28
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin version 3.2
SOFTWARE: Patentin version 3.2
LENGTH: 5
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Pred. No. 1.3e+06;
0; Mismatches 1; Indels
                                                                                                             LOCATION: (2). (2)
OTHER INFORMATION: X can be either F or V
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Publication No. US20040171093A1
GENERAL INFORMATION:
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Best Local Similarity 80.0%;
Matches 4; Conservative
                                                                                                                                                                         FEATURE:

NAME/KEY: MOD RES

LOCATION: (5)...(5)

OTHER INFORMATION: AMIDATION

US-10-695-536-4
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LOCATION: (5)...(5)
CTHER INFORMATION: AMIDATION
US-10-805-881.1
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                                                 FEATURE:
NAME/KEY: MISC_FEATURE
                ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Homo sapiens
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Best Local Similarity
Matches 4; Conserv
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US-10-805-881-1
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US-10-805-881-4
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Sequence 4, Application US/1069536;
Sequence 6, Application No. US20040110692A1
GENERAL INFORMATION:
APPLICANT: Wells, IDEAT Clifton
TITLE OF INVENTION: Antagonists of the Magnesium Binding Defect as Therapeutic Agents;
TITLE OF INVENTION: Antagonists of Treatment of Abnormal Physiological States;
TITLE OF INVENTION: And Methods for Treatment of Abnormal Physiological States;
FILE REFERENCE: 800812-0008
CURRENT APPLICATION NUMBER: US/10/695,536
CURRENT FILING DATE: 2003-10-28
FRIOR APPLICATION NUMBER: US 09/635,266
FRIOR FILING DATE: 2000-08-09
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PatentIn version 3.2
SEQ ID NO 4
ILBRICH: Batter Company of the Company of 
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Sequence 3, Application US/10688741

Sequence 3, Application US/10688741

Sequence 3, Application No. US20040106636A1

SEMENAL INFORMATION:

APPLICANT: Kream, Richard M.

APPLICANT: Kream, Richard M.

APPLICANT: Kream, Richard M.

TITLE OF INVENTION: Method Of Inhibiting Opioid Tolerance Development With Chimeric F.

TITLE OF INVENTION: Analgesics

TITLE OF INVENTION: Analgesics

TITLE OF INVENTION NUMBER: US/10/688,741

CURRENT APPLICATION NUMBER: 2003-10-17

NUMBER OF SEQ ID NOS: 3

SEQ ID NO 3

SEQ ID NO 3
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Pred. No. 1.3e+06;
0; Mismatches 1; Indels
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80.0%; Pred. No. 1.3e+06;
rative 0; Mismatches 1; Indels
APPLICANT: Kream, Richard M.
APPLICANT: Kream, Richard M.
TITLE OF INVENTION: Chimeric Hybrid Analgesics
FILE REPERENCE: Kream
CURRENT APPLICATION NUMBER: US/10/134,187
CURRENT FILING DATE: 2002-04-26
NUMBER OF SEQ ID NOS: 3
SOFTWARE: Patentin version 3.1
SEQ ID NO 3
LENGTH: 5
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80.0%;
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Best Local Similarity 80.07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                     ; TYPE: PRT
; ORGANISM: mammalian
US-10-134-187-3
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; TYPE: PRT
; ORGANISM: mammalian
US-10-688-741-3
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US-10-695-536-4
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APPLICANT: Teruo Nishida
APPLICANT: Teruo Nishida
APPLICANT: Makoto Inui
APPLICANT: Masatsugu Nakamura
TITLE OF INVANTION: NOVEL PEPTIDE AND PHARMACEUTICAL USE OF THE SAME
FILE REFRENCE: 04355/HG
CURRENT APPLICATION NUMBER: US/10/497,628
CURRENT FILING DATE: 2004-06-03
PRIOR PPLICATICATION NUMBER: JP 2001-368103
PRIOR FILING DATE: 2001-12-01
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 30, Application US/08484409;
Publication No. US20020076412A1
GENERAL INFORMATION:
APPLICANT: Steinman, Lawrence
APPLICANT: Zamvil, Scott
TITLE OF INVENTION: METHODS FOR MODULATING THE IMMUNE SYSTEM
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STAFE: Washington
CITY: Seatile
STAFE: Washington
CONNTRY: USA
ZIP: 98104-7092
COMPUTER: IBM PC compatible
GOMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPANION
COMPUTER: IBM PC COMPANION
COMPUTER: IBM PC COMPANION
COMPUTER: IBM PC COMPANION
COMPANION SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    68.0%; Score 17; DB 14; Length 5; 60.0%; Pred. No. 1.3e+06; 1; Indels
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60.0%; Pred. No. 1.3e+06;
iive 1; Mismatches 1
                                                                                                                                                                                                                                       OTHER INFORMATION: Artificial Polypeptide
                                                                                                                                                                                                                                                                                                       ) NAME/KEY: PEPTIDE

) LOCATION: (02)...(02)

: OTHER INFORMATION: Xaa is any amino acid

US-10-168-789A-32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-497-628-16; Sequence 16, Application US/10497628; Publication No. US20050009752A1; GENERAL INFORMATION:
                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 60.0
Matches 3; Conservative
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Best Local Similarity 60.0
Matches 3; Conservative
                                                     NUMBER OF SEQ ID NOS: 64
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-497-628-16
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                                                                                SEQ ID NO 32
LENGTH: 5
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LENGTH: 5
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                                                                                                                                                                                                                                                                        FEATURE:
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APPLICANT: KITADA, Chieko
APPLICANT: KITADA, Chieko
APPLICANT: KITADA, Chieko
APPLICANT: KITADA, Solosoo (1820030148943Aluhiro
TITLE OF INVENTION: No. US20030148943Aluhiro
CURRENT APPLICANTON NUMBER: US/10/168, 789A
CURRENT APPLICANTON NUMBER: US/1000/09083
PRIOR APPLICANTON NUMBER: UP 11-36288
PRIOR FILING DATE: 1999-12-21
PRIOR FILING DATE: 1999-03-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ·.
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| Publication NO US20050009752A1 |
| GENERAL INFORMATION: | APPLICANT: Teruc Nishida |
| APPLICANT: Makero Inui |
| APPLICANT: APPLICANTON NUMBER: JO 040-1368103 |
| BRIOR PELING DATE: 2001-12-01 |
| NUMBER OF SEQ ID NOS: 17 |
| SOFTWARE PATENT OF SEQ ID NOS: 17 |
| SOFTWARE PATENT OF SEQ ID NOS: 17 |
| SOFTWARE PATENT OF SEQ ID NOS: 17 |
| SOFTWARE PATENT OF SEQ ID NOS: 17 |
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Pred. No. 1.3e+06;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Indels
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Pred. No. 1.38+06;
0; Mismatches 1; Indels
                                                                                                                                                                                                            LOCATION: (2)...(2)
OTHER INFORMATION: "X" may be either F or V
FEATURE:
LOCATION: (5)...(5)
OTHER INFORMATION: AMIDATION
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Sequence 32, Application US/10168789A
Publication No. US20030148943A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           80.0%;
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Best Local Similarity 80.0
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 80.0
Matches 4; Conservative
                                                                                TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                            FEATURE:
NAME/KEY: MISC FEATURE
LOCATION: (2). (2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 FVGLM 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 11
US-10-497-628-15
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SEQ ID NO 4
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Sequence 2, Application US/10033026
; Sequence 2, Application US/10033026
; Publication No. US20020147309A1
; GRNERAL INFORMATION:
; GRNERAL INFORMATION:
; APPLICANT: Lipscombe, Diane
; APPLICANT: Lipscombe, Diane
; APPLICANT: Lipscombe, Diane
; TITLE OF INVENTION: HUMAN N-TYPE CALCIUM CHANNEL ISOFORM AND USES THEREOF
; TILLE OF INVENTION: HUMAN N-TYPE CALCIUM CHANNEL ISOFORM AND USES THEREOF
; TILLE OF INVENTION: HUMBER: US/10/033,026
; CURRENT FILING DATE: 2001-12-28
; PRIOR PELICATION NUMBER: US 60/077,901
; PRIOR FILING DATE: 1999-03-13
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 2
; LENGTH: 4
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'o. 1.3e+06; Indels
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SOFTWARE: PatentIn Release #1.0, Version #1.30
                CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/484,409
FILING DATE: 07-JUN-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: MAKI, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 690068.409C1
TELEPHONE: (206) 622-4900
TELEPHONE: (206) 622-4900
TELEPHONE: (206) 622-631
INFORMATION FOR SEQ ID NO: 30: SEQUENCE CHARACTERISTICS:
LEMOTH: 4 minno acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.0%; Pr
Matches 3; Conservative 0;
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; ORGANISM: Homo sapiens
US-10-033-026-2
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Search completed: March 23, 2005, 15:07:06 Job time: 93 secs

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Query marcn
Best Local Similarity
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                                                                                                                     March 23, 2005, 14:37:21; Search time 30 Seconds (without alignments) 12.442 Million cell updates/sec
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/cgn2_6/ptodata/1/iaa/Aa_COMB.pep:*
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/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-07-753-909B-3
US-07-334-553-2
US-08-269-28B-1
US-08-391-910-1
US-08-391-910-1
US-08-391-910-1
US-08-391-910-1
US-08-341-591-61
US-08-441-591-61
US-08-441-135-1
US-08-444-135-1
US-08-444-135-1
US-08-257-966-1
US-08-257-968-1
US-08-257-968-1
US-08-153-847-1
US-08-153-847-1
US-08-153-847-1
US-08-127-904-2
US-08-127-904-12
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                                                                                                                                                                                                                                                                                                                                                              513545 segs, 74649064 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUMMARIES
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Maximum Match 100%
Listing first 45 summaries
                                                                               protein search, using sw model
                                                                                                                                                                                                                                                                                                                      Gapop 10.0 , Gapext 0.5
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                   Copyright
                                                                                                                                                                                                            SEQ2
25
1 fvglm 5
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Match
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                                                                                                                                                                                                              Title:
Perfect score:
                                                                                                                                                                                                                                                                                                  Scoring table:
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                                                                               OM protein -
                                                                                                                                                                                                                                                          Sequence:
                                                                                                                                                                                                                                                                                                                                                              Searched:
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                                                                                                                            Run on:
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No.
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Sequence 33, App Sequence 35, App Sequence 2, Appl Sequence 12, Appl Sequence 1, App Sequence 2, Appl Sequence 2, Appl Sequence 8, Appl Sequence 2, Appl Sequence 2, Appl Sequence 2, Appl Sequence 8, Appl Sequence 8, Appl Sequence 8, Appl Sequence 2, Appl Sequence 2, Appl Sequence 2, Appl Sequence 3, Appl Sequence 3, Appl Sequence 2, Appl Sequence 3, Appl Sequence 3	Family ase	, č
US-09-264-709A-33 US-09-264-709A-35 US-09-264-709A-35 PCT-US94-10475-2 PCT-US94-10475-12 US-09-284A-2 US-09-264-709A-27 US-09-264-709A-27 US-09-264-709A-27 US-09-608-892-16 US-09-608-892-16 US-09-608-892-16 US-09-638-10475-1 US-08-127-904-8 US-08-303-362A-63 US-08-303-362A-63 US-08-303-362A-63 US-08-303-362A-63 US-08-070-301-8 US-09-635-266-3	ALIGNMENTS 198 198 100 100 100 100 100 100	Score 20; DB 1; Length Pred. No. 4.1e+05;
44444000000000444444 666666666666666666	7753 100	80.0%;
6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6	TLT 1 17-753-909B-3 17-753-909B-3 17-753-909B-3 17-151-100-2 18-15-10-3 18-15-10-3 18-15-10-3 18-15-10-3 18-15-10-3 18-15-10-3 18-15-10-3 18-15-10-3 18-15-10-3 18-15-10-3 18-15-10-3 18-10-3	B-3 similarity
229 3320 3320 3321 3321 3324 3324 3324 3324 3324 3324	RESULT 1 US-07-733-909B-3 Sequence 3, Applicat. Sequence 3, Applicat. Sequence 3, Applicat. APPLICANT: Vaudr. APPLICANT: Conjon TITLE OF INVENTION NUMBER OF SEQUENCIONESCONDENCE ADD. STREET: 804 Gr. CONPUTER: IOWA CONPUTER: IOWA CONPUTER: IDWA CONPUTER: IBM OPERATING SYSTE SOFTWARE: Pate: CURSIPICATION NUM FILING DATE: 1 APPLICATION NUM TELEPHONE: (515) INFORMATION FOR SEQUENCE CHARACTEE LENGTH: S amin TELEPHONE: (515) INFORMATION FOR SEQUENCE CHARACTEE LENGTH: S amin TELEPHONE: TYPE: AMINO AC STRANDEDNESS: TOPOLOGY: line MOLECULE TYPE: D TYPE: AMINO AC STRANDEDNESS: CORLGINAL SOURCE: ORGANISM: RANA DEVELOPMENTAL S TISSUE TYPE: D	3-07-753-909E Query Match Best Local S
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US-08-269-288-1
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Matches
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         Gaps
                                                                                                                  Sequence 2, Application US/07934553
Fatent No. 5314690
GENERAL INFORMATION:
APPLICANT: PATTERSON, ROY
APPLICANT: HARRIS, KATHLEEN E
ITTLE OF INVENTION: ANTIBODIES TO SPECIFIC ALLERGENS
ITLE OF INVENTION: ANTIBODIES TO SPECIFIC ALLERGENS
NUMBER OF SEQUENCES:
ADDRESSE: TILTON, FALLON, LUNGWUS & CHESTNUT
STREET: 100 SOUTH WACKER DRIVE
CITY: CHICAGO
STATE: ILLINOIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | Sequence 1, Application US/08269288 |
| Sequence 1, Application US/08269288 |
| Patent No. 5491140 |
| GENERAL INPORMATION: APPLICANT: Behlert, Donald R. APPLICANT: Gehlert, Donald R. APPLICANT: Lunn, William H.W. |
| TILLE OF INVENTION: William H.W. |
| TILLE OF INVENTION: WAPHTHYL TACHYKININ RECEPTOR ANTAGONISTS |
| WUMBER OF SEQUENCES: 4 |
| CORRESPONDENCE ADDRESS: |
| ADDRESSEE: Bli Lilly and Company
     1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Indels
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                                                                                                                                                                                                                                                                                        4.1e+05;
    0; Mismatches
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80.0%; Pred. No. 4.1e
tive 0; Mismatches
 4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 5 amino acids
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  unknown
                                                  FXGLM 5
                          1 FVGLM 5
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                                                                                                      US-07-934-553-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
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Matches
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FILING DATE:
CLASSIFICATION: 530
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                                      -08-418-994-1
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Patent No. 5563133

GENERAL INFORMATION:
APPLICANT: Hipschind, Philip A.
ITLE OF INVENTION: HEXAMETHYLENEIMINYL TACHYKININ RECEPTOR
ITLE OF INVENTION: ANTAGONISTS
NUMBER OF SEQUENCES: 4

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             i, Indels
                                                                                                                                                                                                                                                                                Length 5;
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: BATCHIN Belease #1.0, Version #1.30
SOCTWARE: PatentIN Belease #1.0, Version #1.30
SOCTWARE: PAPLICATION DATA:
APPLICATION NUMBER: US/08/391,910
FILING DATE:
CLASSIFICATION:
NAME: GAPLO, Paul J
NEGISTRATION NUMBER: 36,808
REFERENCE/DOCKET NUMBER: X-9979
                                                                                                                                                                                                                                                                           80.0%; Score 20; DB 1; I
80.0%; Pred. No. 4.1e+05;
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Pred. No. 4.1e+05;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                    NU 9033-CIP2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Eli Lilly and Company
STREET: Lilly Corporate Center
CITY: Indianapolis
STATE: Indiana
REGISTRATION NUMBER: 16,926
REFERENCE/DOCKET NUMBER: NU 9(
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312)-456-8000
FILEPRAX: (312)-456-7776
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: (317) 276-0756
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 80.0%;
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (317) 276-3861
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                   4; Conservative
                                                                                                                                                                                                                    MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   single
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Best Local Similarity
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| Seguence 1, Application U8/08418994
| Patent No. 265566
| GREERAL INFORMATION: | APPLICANT: | Higherind, Philip A. |
| APPLICANT: | Higherind, P
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ZIP: 80111
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 MG storage COMPUTER: IBM compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION NUMBER: US/08/441,591
FILING DATE:
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/391,814
FILING DATE:
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: GAylo, Paul J.
REGISTRATION NUMBER: 36,808
REFERENCE/DOCKET NUMBER: 36,808
REFERENCE/DOCKET NUMBER: 36,008
TELECOMMUNICATION INFORMATION:
TELECHONE: (317) 276-0756
TELEPAK: (317) 276-0756
TELEPAK: (317) 276-3861
INPORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 61, Application US/08441591
Patent No. 5637682
GENERAL INFORMATION:
APPLICANT: NIEUWLANDT, D., GOLD, L. AND WECKER, M.
TITLE OF INVENTION: HIGH-AFFINITY
TITLE OF INVENTION: OLIGONUCLEOTIDE LIGANDS
TITLE OF INVENTION: SUBSTANCE P
NUMBER OF SEQUENCES: 66
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4.1e+05;
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80.0%; Pred. No. 4.1e+0!
Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: Swanson & Bratschun, D.L.C. STREET: 8400 E. Prentice Avenue, Suite 200 STATE: Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/303,362
FILING DATE: 9-SEPTEMBER-1994
PRIOR APPLICATION NUMBER: 07/714,131
FILING DATE: 10-UNE-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/931,473
FILING DATE: 17-AUGUST-1992
PRIOR APPLICATION NUMBER: 07/931,473
FILING DATE: 17-AUGUST-1992
PRIOR APPLICATION NUMBER: 07/931,473
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 08/117,991
FILING DATE: 8-SEPTEMBER 1993
PAPOR APPLICATION DATA:
APPLICATION NUMBER: 07/536,428
FILING DATE: 11-JUNE-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 80.0
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                          / MOLECULE TYPE: peptide
US-08-391-814-1
                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 FVGLM 5
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US-08-441-591-61
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PRIOR APPLICATION NUMBER: 07/564,624
ATTOGRAMY AGENT :12000ANTON: 1000ANTON: 1000ANTON:
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LILLER, Bruce D.

JAPPLICANT: Hipskind, Philip A.

APPLICANT: Howbert, Jeffry J.

APPLICANT: Howbert, Joseph H.

APPLICANT: Lobb, Karen L.

APPLICANT: Muchl, Brian S.

APPLICANT: Mixon, James M.

TITLE OF INVENTION: ANTAGONISTS

NUMBER OF SEQUENCES:

CORRESPONDENCE ADDRESS:

ADDRESSEE: Eli Lilly and Company

STREET: Lilly Corporate Center/Parent

CITY: Indianapolis

STATE: IN

COUNTPV.
                                                                                                         Score 20; DB 1; Length 5;
Pred. No. 4.1e+05;
0; Mismatches 1; Indels
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PAPLICATION NUMBER: US/08/463,874
FILING DATE: 05-UW-1995
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 46285
COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                            US-08-463-874-1
; Sequence 1, Application US/08463874
; Patent No. 5684033
; GENERAL INFORMATION:
APPLICANT: Cho, Sung Y.
; APPLICANT: Crowell, Thomas A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Gaylo, Paul J.
REGISTRATION NUMBER: 36,808
REFERENCE/DOCKET NUMBER: X884
TELECOMMUNICATION INFORMATION:
TELEPHONE: 317-276-0756
                                                                                                                 80.0%;
80.0%;
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SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 80.0°
                                                                                                Query Match
Best Local Similarity 80.0°
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: single
                                  TOPOLOGY: linear

MOLECULE TYPE: peptide

US-08-462-415-1
                      single
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TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
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                  STRANDEDNESS:
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APPLICANT: Crowell, Thomas A.
APPLICANT: Gitter, Bruce D.
APPLICANT: Hipskind, Philip A.
APPLICANT: Howbert, Jeffry J.
APPLICANT: Krushinski, Joseph H.
APPLICANT: Muchl, Karen L.
APPLICANT: Muchl, Brian S.
APPLICANT: Nixon, James A.
TITLE OF INVENTION: HETEROCYCLIC TACHXKININ RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                         80.0%; Score 20; DB 1; Length 5;
80.0%; Pred. No. 4.1e+05;
ive 0; Mismatches 1; Indels
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APPLICATION NUMBER: US/08/462,415 FILING DATE: 05-JUN-1995 CLASSIFICATION: 514 ATORNEY/AGENT INFORMATION: MAME: GAYLO, Paul J. REGITATATION NUMBER: 36,808 REFERENCE/DOCKET NUMBER: X8849B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Eli Lilly and Company
STREET: Lilly Corporate Center/Patent Division
CITY: Indianapolis
                                                                                                                                                                                                                                                                                                                           AROMATIC OR ALIPHATIC AMINO ACID
                                    NAME: Barry J. Swanson
REGISTRATION NUMBER: 33,215
REFERENCE/DOCKET NUMBER: NEX21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 793-3333
INFORMATION FOR SEQ ID NO: 61:
SEQUENCE CHARACTERISTICS:
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5-08-462-415-1
Sequence 1, Application US/08462415
Patent No. 5670499
FILING DATE: 21-OCTOBER-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 317-276-0756
                  ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    4; Conservative
                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Eli Lilly
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                        LOCATION: 2
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                    TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                            NAME/KEY: Xaa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 FVGLM S
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| Patent No. 574482
| GENERAL INFORMATION:
| APPLICANT: Cohen, Marlene L. |
| APPLICANT: Johnson, Kirk W. |
| APPLICANT: Johnson, Kirk W. |
| TITLE OF INVENTION: USE. OF A SEROTONIN AGONIST IN THE TITLE OF INVENTION: COMBINATION WITH A TACHYKININ RECEPTOR ANTAGONIST IN THE TITLE OF INVENTION: TREATMENT OR PREVENTION OF MIGRAINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                        Sequence 1, Application US/08444135
Patent No. 5723575
Patent No. 5723575
Patent No. 5723575
Patent No. 5723576
Patent No. 61100, Chaim
APPLICANT: Glion, Chaim
APPLICANT: Glion, Chaim
APPLICANT: Glioner, Zvi
APPLICANT: Byk, Gerardo
TITLE OF INVENTION: Their Preparation and Pharmaceutical Compositions
TITLE OF INVENTION: Containing Them
NUMBER OF SEQUENCES: 2
CORRESPONDENCE 2
CORRESPONDENCE 2
CORRESPONDENCE 2
ADDRESSE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC Compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/444,135
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          80.0%; Score 20; DB 1; I
80.0%; Pred. No. 4.1e+05;
cive 0; Mismatches 1;
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OTHER INFORMATION: /label= Xaa
OTHER INFORMATION: /note= "Xaa = Phe or Val"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILLING DATE:

CLASSIFICATION: 530

PRIOR APPLICATION DATA:

PELLING DATE: 01-0CT-1992

ATTORNEY AGENT INFORMATION:

NAME: JARKOVSKY, 188aC

REGISTRATION NUMBER: 7754-003-999

TELECOMMUNICATION INFORMATION:

TELEPHONE: 212 869-8864/9741

INFORMATION FOR SEQ ID NO: 1:

INFORMATION FOR SEQ ID NO: 1:

TELEPHONE: 212 869-8864/9741

INFORMATION FOR SEQ ID NO: 1:

SEQUENCE CHARACTERISTICS:

**CHARACTERISTICS: 5 AND ACIDS

**CHARACTERISTICS: 5 AND A
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Best Local Similarity 80.00,
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STRANDEDNESS: single
TOPOLOGY: linear
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MOLECULE TYPE: peptide
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US-08-444-135-1
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US-08-318-391-1
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80.0%; Score 20; DB 1; Length 5;
Best Local Similarity 80.0%; Pred. No. 4.1e+05;
Matches 4; Conservative 0; Mismatches 1; Indels
            CORRESPONDENCE ADDRESS:

ADDRESSEE: Eli Lilly and Company
STREET: Lilly Corporate Center
GITY: Indianapolis
STATE: Indianapolis
STATE: Indianapolis
CONTRY: United States of America
ZIP: 46285
COMPUTER: Eloppy disk
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Ploppy disk
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
COMPUTER: PatentIn Release #1.0, Version #1.25
SOFTWARE: PatentIn Release #1.0, Version #1.25
COMPUTER: DATE: US/08/318,391
FILING DATE: US/08/318,391
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ZIP: 02110-2804

ZIP: 02110-2804

COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compartible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/737,371E
FILING DATE: 29-JUL-1991
CLASSIFICATION DATA:
APPLICATION NUMBER: 07/559,172
APPLICATION NUMBER: 07/559,172
APPLICATION NUMBER: 07/559,172
ATTORNEY AGENT INFORMATION:
NAME: Freeman, John W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Freeman, John W.
REGISTRATION NUMBER: 29,066
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NUMBER OF SEQUENCES:
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Fatent No. 6175013
GENERAL INFORMATION:
APPLICANT: Howbert, James J.
APPLICANT: Muchl, Brian S.
TILLE OF INVENTION: IMIDAZOLINYL TACHYKININ RECEPTOR TITLE OF INVENTION: ANTAGONISTS
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Eli Lilly and Company STREET: Lilly Corporate Center/1104
CITY: Indianapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 80.0%; Score 20; DB 3; Length 5; Best Local Similarity 80.0%; Pred. No. 4.1e+05; Matches 4; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                      Length 5;
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COUNTRY: United States of America
COMPUTER: Tology disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/257,966
FILING DATE:
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: GAYLO, Paul J.
REGISTRATION NUMBER: 36,808
REFERENCE/DOCKET NUMBER: 36,808
REGISTRATION NUMBER: 36,808
REJERAX: (317) 276-0766
ITELEPAX: (317) 276-3861
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: S amino acids
LENGTH: S amino acids
                                                                                                                                                                                                                                                                                                                                 Query Match 80.0%; Score 20; DB 2; I Best Local Similarity 80.0%; Pred. No. 4.1e+05; Matches 4; Conservative 0; Mismatches 1;
REFERENCE/DOCKET NUMBER: 00108/028002
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-8006
TELEFAX: 200154
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                            , MOLECULE TYPE: protein US-07-737-371E-6
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STRANDEDNESS: single
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MOLECULE TYPE: peptide
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US-08-257-966-1
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1 FXGLM 5

Search completed: March 23, 2005, 14:50:59
Job time : 31 secs

GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

March 23, 2005, 15:03:13 ; Search time 38 Seconds
 (without alignments)
 10.128 Million cell updates/sec

Run on:

SEQ3 Title: Perfect score:

21 1 fglm 4 Sequence: **BLOSUM62** Scoring table:

Gapop 10.0 , Gapext 0.5

283416 segs, 96216763 residues Searched: Total number of hits satisfying chosen parameters:

86

Minimum DB seq length: 0 Maximum DB seq length: 4

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

PIR 79:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	scripti	Ig heavy chain CRD	T-cell receptor be		spinal cord peptid	эвота1	T-cell receptor be	Н	_	blood cell protein	-modu	bursin - chicken	spinal cord peptid	Famid	hypothetical prote		hypothetical prote		autho-RF amide neu	RPCH-related neuro	myosin-light-chain	Ig mu chain V regi	ubiquitin - rat	globulin IV alpha	carbon-monoxide de	neuropeptide Antho	neuropeptide Antho	FMRFamide - polych	natin-I -	Ig heavy chain CRD
	ΙD	PT0240	A53284	PT0633	B23751	E44823	B53284	PT0636	PT0571	S68328	GKHU	A60898	A23751	ECXAA	D41654	S53508	T30569	138888	A25844	A34626	839390	843959	847552	S09478	PL0140	JQ1273	A35779	A60418	A32480	PT0271
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ф,	Query Match	47.6	47.6	38.1		33.3	33.3	28.6	28.6	28.6	28.6	28.6	28.6	28.6	28.6	28.6	8		28.6	8	θ.	8	28.6	28.6	28.6	28.6	28.6	28.6	28.6	28.6
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T-cell receptor be	tyrosine-melanocyt	cardioexcitatory n	angiotensin-conver	histidinol dehydro	gene p20K protein	T-cell receptor be	tyrosine protein k	cytochrome-c oxida	thyroglobulin - do							
PT0711	PT0698	PT0677	PT0706	PT0675	PT0721	PT0566	A32039	ECNK	PQ0010	S13894	I50412	PT0578	178890	T13892	S18401	
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30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	

ALIGNMENTS

	RESULT 1 PT0240
_	Ig heavy chain CRD3 region (clone 2-100B) - human (fragment)
	C.) Decision Supramo Carrellian, Commun. C.) Decision 30-Sep-1993 #text_change 16-Aug-1996 C.) Angerian Decision. Decision
	R. Yamada, W. Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
	O. DAP. Med. 179, 353-177, 1331 A;Thle: Preferential utilization of specific immunoglobulin heavy chain diversity and jo A:Reference number: PT022: MITD:91108347: PMTD:1899102
	A; Accession: PT0240 A; Molecule type: DNA
	A;Residues: 1-4 <yam> A:Experimental source: B lymphocyte</yam>
	C; Keywords: heterotetramer; immunoglobulin
	Query Match 47.6%; Score 10; DB 2; Length 4; Best Local Similarity 100.0%: Pred. No. 2.8e+05;
	Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Gaps ö Indels ô Mismatches ö Conservative .. 7 Matches

2 GL 3 5 5 à g

T.-cell receptor beta 2 chain D region, Dbeta2 - rabbit
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 02-May-1994 #sequence_revision 18-Nov-1994 #text_change 05-Nov-1999
C;Date: 02-May-1994 #sequence_revision 18-Nov-1994 #text_change 05-Nov-1999
C;Accession: A52284
R;Harindranath, N.; Alexander, C.B.; Mage, R.G.
Mol. Immunol. 28, 881-888, 1991
A,Title: Evolutionarily conserved organization and sequences of germline diversity and jc
A,Reference number: A52284
A,Recession: A52284
A,Rocession: A52284
A,Rolecule type: DNA
A,Rolecule type: DNA
A,Rolecule type: DNA
A,Rolecule cype: Conserved from NCBI backbone (NCBIN:60737, NCBIP:60739)
C;Keywords: T-cell receptor

Query Match

47.6%; Score 10; DB 2; Length 4;
Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 2; Conservative 0; Mismatches 0; Indels m 2 GL ò

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Gaps

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T-cell receptor beta 2 chain D region, Dbeta2 - rabbit
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Accession: B3284
R;Harindranath, N.; Alexander, C.B.; Mage, R.G.
Mol. Immunol. 28, 881-888, 1991
A;Title: Evolutionarily conserved organization and sequences of germline diversity and jc.
A;Title: Evolutionarily conserved organization and sequences of germline diversity and jc.
A;Reference number: A53284; MUD:91342695; PMID:1678859
A;Recession: B53284
A;Residues: 1-4 - HAR>
A;Residues: 1-4 - HAR>
A;Residues: 1-4 - HAR>
A;Residues: GB:860737; NID:g233916; PIDN:AAB19518.1; PID:g233918
A;Residues: requence extracted from NCBI backbone (NCBIN:60737, NCBIP:60738)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TreelI receptor beta chain V-D-J region (141-1CM) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 15-Jun-2001 #sequence_revision 15-Jun-2001 #text_change 15-Jun-2001
C;Accession: PT051
F;FenerY, A.J.
J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  T-cell receptor beta chain V-D-J region (100-2AT) - mouse (fragment)
C;Species: Mns musculus (house mouse)
C;Date: 15-Jun-2001 #sequence_revision 15-Jun-2001 #text_change 15-Jun-2001
C;Accession: PT0636
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50.0%; Pred. No. 2.8e+05;
Live 1; Mismatches 0; Indels
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A;Molecule type: mRNA
A;Residues: 1-3 <FBE>
A;Experimental source: day 19 fetal thymus, strain BALB/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Experimental source: newborn thymus, strain BALB/c C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Status: translation not shown
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Best Local Similarity
Matches 1; Conserv
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                                                                     C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 09-Jul-2004
C;Accession: PT0633
R;Feersion: PT0633
J; Exp., A.J.
J; Exp., A.J.
J; Exp., Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0639; MUID:91277601; PMID:1711558
A;Accession: PT0633
A;Accession: PT0633
A;Accession: PT0633
A;Accession: PT0633
A;Accession: PT0633
A;Cross_references: UNIPROT:Q8BIV7
A;Experimental source: newborn thymus, strain BALB/C
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Windprosomal-associated protein SNAP-25 peptide 1 - rabbit (fragment)
NiAlternate names: superprotein peptide 1
Cispecial Oryctolagus cuniculus (domestic rabbit)
Cispecial Substance Sequence revision 18-Nov-1994 #text_change 15-Jun-1996
Ciscession: E4483
Riboew, A.; Liu, W.S.; Baitinger, C.; Willard, M.B.
Neurosci. 11, 3412-3421, 1991
Airile: The major 35S-methionine-labeled rapidly transported protein (superprotein) is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               pinal cord peptide SCP-5 - pig
Cispecies: Sus scrofa domestica (domestic pig)
Cispecies: Sus scrofa domestica (domestic pig)
Cispecies: 5-Jun-2001 #sequence_revision 15-Jun-2001 #text_change 15-Mar-2004
Cispecies: 5-Jun-2001 #sequence_revision 15-Jun-2001 #text_change 15-Mar-2004
Sishsi, K.L.; Chen, R.L.; Chen, Z.G.; Zhang, H.L.; Lu, Y.A.; Guo, S.Y.; Wu, S.X.; Tsou, Areference number: Biophys. 240, 178-183, 1985
AiReference number: A23751; MUID:85250425; PMID:4015098
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                        -cell receptor beta chain V-D-J region (120-2C) - mouse (fragment)
Species: Mus musculus (house mouse)
19ate: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 09-Jul-2004
Accession: PT0633
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Pred. No. 2.8e+05;
1; Mismatches 0; Indels
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Pred. No. 2.8e+05;
1; Mismatches 0; Indels
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A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-4 <1.OE
A;Experimental source: visual tissue
A;Note: sequence extracted from NCBI backbone (NCBIP:64247)
C;Keywords: membrane trafficking
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Best Local Similarity 50.0%;
Matches 1; Conservative ;
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A; Molecule type: protein
A; Residues: 1-3 <HSI>
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A;Accession: A01421
A;Molecule type: protein
A;Rolecilues: 1-3 <SCH>
A;Note: this serum tripeptide is found to stimulate growth of some cell types and to inh
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C;Species: Gallus gallus (chicken)
C;Date: 15-Jun-2001 #sequence_revision 15-Jun-2001 #text_change 15-Mar-2004
C;Accession: A60898
C;Accession: A60898
R;Audhya, T.; Kroon, D.; Heavner, G.; Viamontes, G.; Goldstein, G.
Science 231, 997-999, 1986
A;Title: Trippetide structure of bursin, a selective B-cell-differentiating hormone of A;Reference number: A60898; MUID:86122916; PMID:3484838
                                                                                                                                                                                                                                                                                                               Diod cell protein A - Molgula manhattensis (fragment)
Cispecies: Molgula manhattensis
Cispecies: Molgula manhattensis
Cispecies: Is-Jun-2001 #sequence_revision 15-Jun-2001 #text_change 15-Jun-2001
Ciscession: S68328
RiTaylor, S.W.; Ross, M.M.; Waite, J.H.
Raylor, S.W.; Ross, M.M.; Waite, J.H.
Arch. Blochem. Blophys-324, 228-240, 1995
A;Title: Novel 3,4-di- and 3,4-5-trihydroxyphenylalanine-containing polypeptides from the A;Reference number: S68325; MUID:96132650; PMID:8554314
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C;Species: Homo sapiens (man)
C;Accession: A01421
R;Schlesinger, D.H.; Pickart, L.; Thaler, M.M.
Experientia 33, 324-325, 1977
A;Title: Growth-modulating serum tripeptide is glycyl-histidyl-lysine.
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A;Residues: 1-3 <AUD>
C;Keywords: amidated carboxyl end; hormone
F;3/Modified site: amidated carboxyl end (Gly) #status experimental
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                                                   Length 3;
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ative 0; Mismatches
                                                                                           1; Conservative
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C; Keywords: T-cell receptor
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spinal cord peptide SCP-4 - pig
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 15-Jun-2001 #sequence_revision 15-Jun-2001 #text_change 15-Mar-2004
C;Accession: A23751
R;Hsi, K.L.; Chen, R.L.; Chen, Z.G.; Zhang, H.L.; Lu, Y.A.; Guo, S.Y.; Wu, S.X.; Tsou, K. Arch. Biochem. Biophys. 240, 178-183, 1985
A;Reference number: A23751; MUID:85250425; PMID:4015098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Accession: A26666
R;Grimmelikhuijzen, C.J.P.; Graff, D.
Proc. Natl. Acad. Sci. U.S.A. 83, 9817-9821, 1986
A;Title: Isolation of <Glu-Gly-Arg-Phe-NH2 (Antho-RFamide), a neuropeptide from sea anemc A;Reference number: A26666; MUID:87092339; PMID:2879288
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A.Residues: 1-4 <GRI>
A.Residues: 1-14 <GRI>
A.Cromsor-eferences: UNIPROT:P10419
G.Comment: The function of this peptide is not known but it could act as a transmitter at C.Comment: Synthetic and natural peptides had identical properties.
G.Superfamily: Refamide neuropeptide
G.Keywords: amidated carboxyl end; neuropeptide; pyroglutamic acid
F.1/Modified site: pyrolidone carboxylic acid (Gln) #status experimental
F.4/Modified site: amidated carboxyl end (Phe) #status experimental
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Da1654
hypothetical protein (sodC 5' region) - Haemophilus parainfluenzae (fragment)
hypothetical protein (sodC 5' region) - Haemophilus parainfluenzae
c)Species: Haemophilus parainfluenzae
c)Species: Haemophilus parainfluenzae
c)Date: 12-Jun-1992 #sequence_revision 12-Jun-1992 #text_change 24-Feb-1995
C)Accession: D41654
B; Kroll, J.S.; Langford, P.R.; Loynds, B.M.
J. Bacteriol. 173, 7449-7457, 1991
A; Title: Copper-zinc superoxide dismutase of Haemophilus influenzae and Haemophilus para:
A; Reference number: A41654; MUID:92041655; PMID:1938942
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C;Species: Anthopleura elegantissima
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A;Molecule type: protein
A;Residues: 1-3 <HSI>
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seq3.rpr
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Starvation-induced ribonuclease - tomato
C.Species: Lycopersicon esculentum (tomato)
C.Species: Lycopersicon esculentum (tomato)
C.Acession: 553508
R.Koeck, M.; Loeffler, A.; Abel, S.; Glund, K.
Plant Mol. Biol. 27, 477-485, 1995
A.Fitle: cDNA structure and regulatory properties of a family of starvation-induced ribc A, Accession: 553508
A, A, Accession: 553508
A, Reference number: 553508
A, Reference preliminary
A, Molecule type: protein
A, Molecule type: protein
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A, Accession: D41654
A, Status: preliminary
A, Molecule type: DNA
A, Residues: 1-4 < KRO>
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Search completed: March 23, 2005, 15:13:39 Job time : 38 secs

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GenCore version 5.1.6 (c) 1993 - 2005 Compugen Ltd.
                     Copyright
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OM protein - protein search, using sw model

Run on:

March 23, 2005, 14:51:07; Search time 171 Seconds (without alignments) 11.978 Million cell updates/sec

SE03 Title: Perfect score:

21 1 fglm 4 Sequence:

BLOSUM62 Scoring table:

Gapop 10.0 , Gapext 0.5

1612378 segs, 512079187 residues Searched:

56 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 4

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

UniProt_03:*
1: uniprot_sprot:*
2: uniprot_trembl:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Description	P58648	P83568	P83570	P01157	P35904	P19916	P02731	P42562	P42563	P58705	P42561	P58707	P01162	P58706	P58649	Q16047	P19918	Q96at0	008433	P24272	P36515	P62970	P62971	P62968	P62969	P01858
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	OCP1 OCTM	E SEPOF	SEPOF	GRWM HUMAN	ACHE	DCML_PSECH	EOSI HUMAN	FAR3 HIRME	FAR4 HIRME	FFKA ANTEL	FLRF HIRME	FLRN ANTEL	FMRF MACN	FYRI_ANTEL	OCP3_OCTM1	47	DCMS PSECH	O.E.	33	VIBE	_YEAST	THYL BOMOR	THYL NOTVI	THYL PIG	THYL_SHEEP	HUMAN
GI .	OCP1	ILME	GWA	GRWM	ACH1	DCML	EOSI	FAR3	FAR4	FFKA	FLRF	FLRN	FMRF	FYRI	OCP3	016047	DCMS	Q96AT0	008433	LUXE	YLM1	THYL	THYL	THYL	THYL	TUFT
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Query Match	57.1	42.9	28.6	28.6	28.6	28.6	28.6	28.6	28.6	28.6	28.6	28.6	28.6	28.6	28.6	28.6	23.8	23.8	19.0	9.5	4.8	0.0	0.0	0.0	0.0	0.0
Score	12	0	9	9	9	9	9	9	9	9	9	9	9	9	9	9	2	S	4	7	r	0	0	0	0	0
Result No.	-1	8	m	4	٠.	9	7	80	6	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	56

ö than Ocp-1.
-- SUBCELLULAR LOCATION: Secreted.
-- PTW. Cop-2 has L-Phe instead of D-Phe.
-- PTW. Cop-2 has L-Phe instead of D-Phe.
-- MASS SPECTROMETRY: MM=395.2; METHOD=MALDI; RANGE=1-4; NOTE=Ref.1.
D-amino acid; Direct protein sequencing; Hormone.
MOD_RES 2 D-phenylalanine (in form Ocp-1).
SEQUENCE 4 AA; 394 MW; GAAR79C8L0000000 CRC64; MEDLINE=20336815; PubMed=10876044; DOI=10.1016/S0196-9781(00)00201-1; Iwakoshi E., Hisada M., Minakata H.;
"Cardioactive peptides isolated from the brain of a Japanese octopus,
Octopus minor."; Gaps Peptides 21:623-630 (2000). -!- FUNCTION: Cardioactive; has both positive chronotropic and inotropic effects on the heart. Ocp-2 is a 1000 time less active 28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
28-FEB-2003 (Rel. 44, Last sequence update)
Cardioactive peptides Ocp-1/Ocp-2.
Octopus minor (Octopus).
Ekkaryota; Metazoa; Mollusca; Cephalopoda; Coleoidea; Neocoleoidea; Octopodiformes, Octopoda; Incirrata; Octopodidae; Octopus.
NISI_TAXID=89766; ö SEQUENCE, SYNTHESIS, MASS SPECTROMETRY, AND CHARACTERIZATION 57.1%; Score 12; DB 1; Length 4; 100.0%; Pred. No. 1.6e+06; tive 0; Mismatches 0; Indels 4 AA. PRT; Query Match Best Local Similarity luv... 2, Conservative STANDARD; 1 FG 2 TISSUE=Brain; OCTMI ઠ 셤

Bukaryota, Metazoa, Mollusca, Cephalopoda, Coleoidea, Neocoleoidea, Decapodiformes, Sepioidea, Sepidae, Sepia. NCBI_TaxID=6610; TISSUB=Egg; MEDLINE=20403899; PubMed=10944467; DOI=10.1006/bbrc.2000.3286; SEQUENCE, SUBCELLULAR LOCATION, TISSUE SPECIFICITY, AND MASS P83568, 29-MAR-2004 (Rel. 43, Created) 29-MAR-2004 (Rel. 43, Last sequence update) 05-JUL-2004 (Rel. 44, Last annotation update) Pheromone peptide ILME. Sepia officinalis (Common cuttlefish). 4 AA. PRT; STANDARD; officinalis."; ILME SEPOF ILME_SEPOF

TISSUE=Egg; MEDLINE=22197108; PubMed=12207899; DOI=10.1016/S0006-291X(02)02036-3; Zatylny C., Gagnon J., Boucaud-Camou E., Henry J.; "ILME: a waterborne pheromonal peptide released by the eggs of Sepia Biochem. Biophys. Res. Commun. 275:217-222(2000). SEQUENCE

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residue."
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SEQUENCE
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ACH1_ACHFU
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DCML_PSECH
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X MEDLINE-98100358; PubMed=9437704; DOI=10.1016/S0196-9781(97)00241-6;

XA Henry J., Favrel P., Boucaud-Camou E.;

RT Testale in and identification of a novel Ala-Pro-Gly-Trp-amide-related cuttlefish, Sepia officinalis.";

R Peptide inhibiting the motility of the mature oviduct in the cuttlefish, Sepia officinalis.";

C --- FUNCTION: Regulatory neuropeptide with myotropic activity

C --- FUNCTION: Regulatory neuropeptide of contractions.

C --- FUNCTION: Regulatory neuropeptide of contractions.

C --- FUNCTION: Secreted.

C --- FUNCTION: Secreted.

--- MASS SPECTROMETRY: MW=259.9; METHOD=MALDI; RANGE=1-2; NOTE=Ref.1.

KW Model RES --- RAS SPECTROMETRY: MW=259.9; METHOD=MALDI; RANGE=1-2; NOTE=Ref.1.

FT MOD RES --- Tryptophan amide.

S SEQÜENCE 2 AA; 261 MW; 73781000000000 CRC64;
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                                                        Gaps
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Eukaryota, Metazoa, Mollusca, Cephalopoda, Coleoidea, Neocoleoidea,
Decapodiformes, Sepioidea, Sepiidae, Sepia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens (Human).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eukerza; Primates; Catarrhini; Hominidae; Homo.
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                       42.9%; Score 9; DB 1; Length 4; 100.0%; Pred. No. 1.6e+06; ive 0; Mismatches 0; Indeli
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                                                                                                                                                                                                                                                                                SEQUENCE, FUNCTION, MASS SPECTROMETRY, AND AMIDATION.
4 AA; 505 MW; 6B1697203000000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.6e+06;
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21-JUL-1986 (Rel. 01, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Growth-modulating peptide.
                                                                                                                                                           2 AA.
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100.0%; Pred. No. 1.6
tive 0; Mismatches
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                                               Conservative
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Best Local Similarity
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Best Local Similarity
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GRWM_HUMAN
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-!- FUNCTION: Neuroexcitatory peptide; increases the impulse frequency
and produces a spike broadening of the identified heart excitatory
neuron (PON); also enhances the amplitude and frequency of the
heart beat. Has also an effect on several other muscles.
PIR; A32480; A32480.
                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Mollusca; Gastropoda; Pulmonata; Stylommatophora;
Sigmurethra; Achatinoidea; Achatinidae; Achatina,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIINE=93014529; PubMed=1399265; Ishida T., In Y., Doi M., Inoue M., Yasuda-Kamatani Y., Minakata H., Ishashita T., Nomoto K., "Crystal structure and molecular conformation of achain-I (H-Gly-D-Phe-Ala-Asp-OH), an endogenous neuropeptide containing a D-amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN-Ferussac; TISSUE-Ganglion;
MEDLINE-8927351; PubMed-2597281;
Kamatani Y., Minakata H., Kenny P.T.M., Iwashita T., Watanabe K.,
Funase K., Sun X.P., Yongairi A., Kim K.H., Novales-Li P.,
Novales E.T., Kanapi C.G., Takeuchi H., Nomoto K.;
"Achatin-I. an endogenous neuroexcitatory terrapeptide from Achatina fulica Ferussac containing a D-amino acid residue.";
Biochem. Biophys. Res. Commun. 160:1015-1020(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN-Ferussac; TISSUE-Heart atrium; MEDLINE=91264856; PubMed=1675568; Fujimoto K., Kubota I., Yasuda-Kamarani Y., Minakata H., Nomoto K., Yoshida M., Harada A., Muneoka Y., Kobayashi M.; "Purification of achatin-I from the atria of the African giant snail, Achatina fulica, and its possible function."; Biochem. Biophys. Res. Commun. 177:847-853 (1991).
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                      DB 1; Length 3; . 1.6e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  D-anino acid; Direct protein sequencing; Hormone.
MOD RES 2 2 D-phenylalanine.
SEQUENCE 4 AA; 408 MW; 6AADD9C810000000 CRC64;
                                                                                                                                                                                                                                                                                                                                                         01-JUN-1994 (Rel. 29, Created)
01-JUN-1994 (Rel. 29, Last sequence update)
25-OCT-2004 (Rel. 45, Last annotation update)
28.6%; Scor.
100.0%; Pred. No. ...
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Achatina fulica (Giant African snail).
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P19916;
01-FEB-1991 (Rel. 17, Created)
                                                                   1; Conservative
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les 1, Conserv
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P42562;
01-NOV-1995 (
01-NOV-1995 (
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-!- MISCELLANEOUS: These peptides are released from mast cells in lung (and other tissues) duting hypersensitivity reactions (anaphylaxis). Their activities, preferentially affecting eosinophils, include chemotaxis, chemotactic deactivation, release of enzymes, and stimulation of the hexose monophosphate shunt.

GO; GO:0006935; P:chemotaxis, IDA.

GO: GO:0006955; P:chemotaxis, IDA.

Hower peptide).
                                                                                                                                                                                                                                           -!- CATALYTIC ACTIVITY: CO + H(2)O + A = CO(2) + AH(2).
-!- COPACTOR: Binds 1 copper(1) ion, 1 molybdenum(VI) ion and 1 molybdopterin cytosine dinucleotide (MCD) per subunit.
-!- SUBDNIT: Heterotrimer consisting of a large, a medium and a small
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Goetzl E.J., Austen K.F.;
"Purification and synthesis of eosinophilotactic tetrapsptides of
human lung tissue: identification as eosinophil chemotactic factor of
anaphylaxis.";
                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                              Kraut M., Hugendieck I., Herwig S., Meyer O.;
"Homology and distribution of CO dehydrogenase structural genes in
                                                                                                                                                                                                                   -!- FUNCTION: Catalyzes the oxidation of carbon monoxide to carbon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
NCBI_TaxID=9606;
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                                                                               Bacteria; Proteobacteria; Alphaproteobacteria; Rhizobiales;
Bradyrhizobiaceae.
                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
01-FEB-1991 (Rel. 17, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Carbon monoxide debydrogenase large chain (EC 1.2.99.2)
dehydrogenase subunit L) (CO-DH L) (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                    Length 4;
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                                                                                                                                                                                                                                                                                                              PIR; PL0140; PL0140.
Direct protein sequencing; Molybdenum; Oxidoreductase.
                                                                                                                                                                                                                                                                                                                                                    SEQUENCE 4 AA; 441 MW; 7761E876F000000 CRC64;
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100.0%; Pred. No. 1.6e+06;
ative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 AA.
                                                                                                                                                                                         carboxydotrophic bacteria.";
Arch. Microbiol. 152:335-341(1989).
                                                                                                                                               MEDLINE=90055678; PubMed=2818128;
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                                                                  Pseudomonas carboxydohydrogena
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21-JUL-1986 (Rel. 01, Last seq
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                                                                                                                                                                                                                                                                                                                                                               MEDLINE=92195954; PubMed=1686933; DOI=10.1016/0196-9781(91)90035-N; Evans B.D., Pohl J., Kartsonis M.A., Calabrese R.L.; "Indentification of Framide neuropeptides in the medicinal leech."; Peptides 12:897-908(1991).
-i. SUBCELLULAR LOCATION: Secreted.
-i. SIMILARITY: Belongs to the FARP (FMRFamide related peptide)
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01-NOV-1995 (Rel. 32, Last sequence update)
05-JUD-2004 (Rel. 44, Last annotation update)
05-JUD-2004 (Rel. 44, Last annotation update)
FMRFamide-like neuropeptide YMRF-amide
Hirudo medicinalis (Medicinal leech).
Eukaryota; Metazoa; Annelida; Clitellata; Hirudinida; Hirudinea;
Arhynchobdellida; Hirudiniformes; Hirudinidae; Hirudinea;
                                                                                                                                         FMRFamide-like neuropeptide YLRF-amide.
Hirudo medicinalis (Medicinal leech).
Eukaryota; Metazoa; Annelida; Clitellata; Hirudinida; Hirudinea;
Arhynchobdellida; Hirudiniformes; Hirudinidae; Hirudo.
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    -!- SIMILARITY: Belongs to the FARP (FMRFamide related peptide)

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MOD RES 4 4 Phenylalanine amide.
SEQUENCE 4 AA; 616 MW; 69D4068B30000000 CRC64;
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4 AA; 598 MW; 69D4073B30000000 CRC64;
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Pred. No. 1.6e+06;
                                                   01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
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100.0%; Pred. No. ...
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SEQUENCE
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FLRN_ANTEL
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                                                                                                                                                                                                                MEDLINE=93391436; PubMed=8397415;
MCFATLANE I.D., Hudman D., Nothacker H.-P., Grimmelikhuijzen C.J.P.;
MTPH expansion behaviour of sea anemones may be coordinated by two inhibitory neuropeptides, Antho-Kamide and Antho-Riamide.";
Proc. R. Soc. Lond. B. Biol. Sci. 253:183-188(1993).
-!-FUNCTION: Inhibits spontaneous contractions in several muscle groups. May be involved in the expansion phase of feeding behaviour in sea anemones.
                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SPECIES=H.trivolvis; TISSUE=Kidney;
MEDLINE=94286417; PubMed=7912428; DOI=10.1016/0196-9781(94)90166-X;
Madrid K.P., Price D.A., Greenberg M.J., Khan H.R., Saleuddin A.S.M.;
"PMRFamide-related peptides from the kidney of the snail, Helisoma
                                                                                                                                      MEDLINE=92028852; PubMed=1681803; Orimmelikhuijzen C.J.P.; Nothacker H.-P., Rinehart K.L. Jr., Grimmelikhuijzen C.J.P.; Isolation of L-3-phenyllactyl-Phe-Lys-Ala-NH2 (Antho-KAamide), a novel neuropeptide from sea anemones "; Biochem. Biophys. Res. Commun. 179:1205-1211(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SPECIES=H.medicinalis;
MEDLINE=22195554; PubMed=1686933; DOI=10.1016/0196-9781(91)90035-N;
Evans B.-D., Pohl J., Kartsonis M.A., Calabrese R.L.;
"Identification of RFamide neuropeptides in the medicinal leech.";
Peptides 12:897-908(1991).
                                                                            Anthopleura elegantissima (Sea anemone).
Eukarytota Metazoa; Cnidaria; Anthozoa; Zoantharia; Actiniaria;
Nynantheae; Actiniidae; Anthopleura.
NCBI_TaxID=6110;
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Arhynchobdellida; Hirudiniformes; Hirudinidae; Hirudo.
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28-FEB-2003 (Rel. 41, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
                  4 AA.
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01-NOV-1995 (Rel. 32, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
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100.0%; Pred. No. 1.6
tive 0; Mismatches
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              STANDARD.
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hes 1; Conservative
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FFKA ANTEL
ID FFKA ANTEL
AC P58705;
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Proc. Natl. Acad. Sci. U.S.A. 87:5410-5414(1990).
Proc. Natl. Acad. Sci. U.S.A. 87:5410-5414(1990).
Proc. Natl. Acad. Sci. U.S.A. 87:5410-5414(1990).
Proc. Natl. Acad. Sci. U.S.A. Neuron specific.
PRSUS SPECTROMETRY: MW=549.3; METHOD=FAB; RANGE=1-4; NOTE=Ref.1.
PRSUS SPECTROMETRY: MW=549.3; METHOD=FAB; RANGE=1-4; NOTE=Ref.1.
PRSUS SPECTROMETRY: MW=549.3; METHOD=FAB; RANGE=1-4; NOTE=Ref.1.
PROC. Addation; Direct protein sequencing; Neuropeptide.
MOD RES
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Nersis virens (Sandworm),
Hirdo medicinalis (Medicinal leech), and
Helisoma trivolvis (Snail)
Bukaryota; Metazca; Mollusca; Bivalvia; Heteroconchia; Veneroidas;
NCBL TaxID=6594, 6353, 6421, 27815;
Peptides 15:31-36(1994).
-!- SUBCELLULAR LOCATION: Secreted.
-!- SINILARITY: Belongs to the FARP (FMRFamide related peptide)
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100.0%; Pred. No. 1.6e+06;
iive 0; Mismatches 0; Indels
                                                                                               Amidation; Direct protein sequencing; Neuropeptide.
MOD RES
SEQUENCE 4 AA; 582 MW; 69D40729A0000000 CRC64;
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21-JUL-1986 (Rel. 01, Last sequence update)
05-JUL-2004-(Rel. 44, Last annotation update)
FMRFamide (Peak C) (Cardioexcitatory neuropeptide).
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28-FEB-2003 (Rel. 41, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
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                                                                                                                                                                                                                                                                     1; Conservative
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SEQUENCE, AND SYNTHESIS.
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Search completed: March 23, 2005, 15:10:03 Job time : 172 sec8
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"Purification and characterization of a cardioexcitatory neuropeptide from the central ganglia of a bivalve mollusc.";
Prep. Blochem. 7:261-281(1977).
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MEDLINE-94286417; Pubmed=791248; DOI=10.1016/0196-9781(94)90166-X;
Madrid K.P., Price D.A., Greenberg M.J., Khan H.R., Saleuddin A.S.M.;
"FWRFamide-related peptides from the kidney of the snail, Helisoma
                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptides 15:31-36(1994).
-1- FUNCTION: Myoactive; cardioexcitatory substance. Pharmacological activities include augmentation, induction, and regularization of cardiac contraction.
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                                                                                                                                                                                                        MEDLINE=90259866; PubMed=2342992; DOI=10.1016/0196-9781(90)90113-J; Krafilak K.G., Price D.A.; "Authentic PMRFamide is present in the polychaete Nereis virens."; Peptides 11:75-77(1990)
                                                                                                                                                                                                                                                                                 SPECIES=H.medicinalis;
MEDLINE=92195954; PubMed=1686933; DOI=10.1016/0196-9781(91)90035-N;
MEDLINE=92195954; PubMed=1686933; DOI=10.1016/0196-9781(91)90035-N;
Faran 8.D. Dobl J., Kartsonis M.A., Calabrese R.L.;
"Identification of Reamide neuropeptides in the medicinal leech.";
Peptides 12:897-908(1991).
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Nothacker H.-P., Rinehart K.L. Jr., McFarlane I.D.,
Grimmelikhuijzen C.J.P.;
SPECIES=M.nimbosa; TISSUE=Cerebral pedal, and Visceral ganglion;
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Eukaryota; Metazoa; Cnidaria; Anthozoa; Zoantharia; Actiniaria;
Nynantheae; Actiniidae; Anthopleura.
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-!- SIMILARITY: Belongs to the FARP (FWRFamide related peptide)
                      Price D.A., Greenberg M.J.; "Structure of a molluscan cardioexcitatory neuropeptide."; Science 197:670-671 (1977).
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Amidation; Direct protein sequencing; Neuropeptide.
MOD RES
SEQUENCE 4 AA; 600 MW; 69D40699A0000000 CRC64;
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28-FEB-2003 (Rel. 41, Last sequence update)
29-UJU-2004 (Rel. 44, Last annotation update)
Antho-Rlamide I [Contains: Antho-Rlamide II].
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                                                                         SEQUENCE, AND CHARACTERIZATION.
SPECIES=M.nimbosa; TISSUE=Ganglion;
MEDLINE=78012038; PubMed=909875;
              MEDLINE=77215956; PubMed=877582;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Conservative
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                                                                                                                                                                               SEQUENCE.
SPECIES=N.virens;
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FYRI ANTEL
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WEDLINE=33391436; PubMed=8397415;

MAFAILANE=33391436; PubMed=8397415;

MCFAILANE=33391436; PubMed=8397415;

MCFAILANE=33391446; PubMed=8397415;

MCFAILANE=3331446; PubMed=8397415;

MCFAILANE=3331446; PubMed=8397415;

MCFAILANE=3331446; PubMed=8397415;

MCFAILANE=3331446; PubMed=8397415;

MCFAILANE=3331446; PubMed=8397446;

MCFAILA
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MEDLINE=20336815; PubMed=10876044; DOI=10.1016/S0196-9781(00)00201-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Iwakoshi E., Hisada M., Minakata H.;
"Cardioactive peptides isolated from the brain of a Japanese octopus,
Octopus minor."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -!- PTM: Ocp-4 has D-Ser instead of L-Ser.
-!- MASS SPECTROMETRY: WM=395.2; METHOD=WALDI; RANGE=1-4; NOTE=Ref.1.
D-amino acid; Direct protein sequencing; Hormone.
2 2 D-serine (in form Ocb-4).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                    unusual,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FUNCTION: Cardioactive; has both positive chronotropic and inotropic effects on the heart. Ocp-4 is a 1000 time less active
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Mollusca; Cephalopoda; Coleoidea; Neocoleoidea; Octopodiformes; Octopoda; Incirrata; Octopodidae; Octopus.
NCBI_TaxID=89766;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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"Isolation of two novel neuropeptides from sea anemones: the biologically active L-3-phenyllactyl-Tyr-Arg-Ile-NH2 and its phenyllactyl fragment Tyr-Arg-Ile-NH2."; Peptides 12:1165-1173(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE, SYNTHESIS, MASS SPECTROMETRY, AND CHARACTERIZATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28.6%; Score 6; DB 1; Length 4; 100.0%; Pred. No. 1.6e+06; ive 0; Mismatches 0; Indele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 4;
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4 AA; 463 MW; 6AB365B81000000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 4 1 180leucine amide.
4 AA; 598 MW; 60441B59A0000000 CRC64;
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Pred. No. 1.6e+06;
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28-FEB-2003 (Rel. 41, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Cardioactive peptides Ocp-3/Ocp-4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28.6%;
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Best Local Similarity
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OM protein - protein search, using sw model

March 23, 2005, 14:52:02; Search time 164 Seconds (without alignments) 9.433 Million cell updates/sec Run on:

SEQ3 Title:

21 1 fglm 4 Perfect score: Sequence:

BLOSUM62 Scoring table:

Gapop 10.0 , Gapext 0.5

2105692 segs, 386760381 residues Searched:

19815 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 4

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

A Geneseq 16Dec04:* 1: qeneseminî. Database

geneseqp2002s:* geneseqp2003as:* geneseqp2003bs:* geneseqp2004s:* geneseqp1980s:* geneseqp1990s:* geneseqp2000s:* geneseqp2001s:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

STIMMADIES

	Description	Aaw41683 Peptide u	Aay31075 Non-cross	Aab23026 Human/rat	Aay67577 P antagon	Aab91447 Tachykini	Abb10091 Substance	Aau77846 Tachykini	Ade94198 High acti	Adr43772 Human mag	Aap61654 Sequence		-	Abb10092 Substance	Aap61707 Sequence				Aag62647 Typical t			Aay67578 Pantagon	Aab91448 Tachykini	Aap60334 Peptide w	-	Aaw41684 Tetrapept
SUMMAKIES	ID	AAW41683	AAY31075	AAB23026	AAY67577	AAB91447	ABB10091	AAU77846	ADE94198	ADR43772	AAP61654	AAP71301	AAW41686	ABB10092	AAP61707	AAP71312	AAY23485	AAB12293	AAG62647	ABB88046	ADL78809	AAY67578	AAB91448	AAP60334	AAW77469	AAW41684
	Length DB	4 2	4 2	4	4 3	4	4 5	4	4 7	4 8	4	4 1	4 2	4	4	4	4	4	4	4	4	3	3 4	4	4 2	4
æ	Query Match I	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	85.7	85.7	85.7	85.7	76.2	76.2	76.2	76.2	76.2	76.2	76.2	71.4	71.4	71.4	71.4	71.4
	Score	21	21	21	21	21	21	21	21	21	18	18	18	18	16	16	16	16	16	16	16	15	15	15	15	15
	Result No.	-	7	٣	4	S	9	7	æ	6	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25

-	-	Aap71306 Peptide c Aar34486 FGIA. 8/1	Aar46020 Serine pr Aar93149 Mycobacte	Aab12292 Prodrug o	Aab91714 Opioid pe		Abg32223 Sheep col	Adq91509 HIV trunc	Ads74433 Ovine col	Abg77484 Targettin		Aap71287 Opiate bi		8	Aar47299 Peptide a
2 AAW41685 4 AAB91795	4 AAB91822 1 AAP61659		2 AAR93149	3 AAB12292	4 AAB91714	5 ABB88045	5 ABG32223	8 ADQ91509	8 ADS74433	5 ABG77484	1 AAP61658	1 AAP71287	1 AAP82691	2 AAR15768	2 AAR47299
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71.4	66.7	66.7 66.7	66.7	66.7	. 66.7	66.7	66.7	66.7	66.7	61.9	61.9	61.9	61.9	61.9	61.9
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25	20 8	30 31	33 33	34	35	36	37	38	39	40	41	42	43	44	45

ALIGNMENTS

RESULT 1

AAW41683 standard; peptide; 4 AA. (first entry) 09-JUN-1998 AAW41683; AAW41683

Peptide used in ophthalmic drug to treat corneal disorders.

Ophthalmic drug; corneal disorder; ulcer; epithelial peeling; dry eye; keratitis; insulin like growth factor-I; IGF-I; eye drop.

Synthetic.

/note= "C-terminal amide" Location/Qualifiers Key Modified-site

31-DEC-1997.

WO9749419-A1

97WO-JP002015. 11-JUN-1997; 96JP-00165612. (SANT) SANTEN PHARM CO LID. 26-JUN-1996;

Nishida T, Nakamura M, Nakata K;

WPI; 1998-076907/07.

Ophthalmic drug composition containing tetra:peptide - is useful as corneal disorder remedy for corneal ulcer, corneal epithelial peeling, dry eye, keratitis.

Claim 1; Page 15; 19pp; Japanese.

The present sequence represents a tetrapeptide which is the active ingredient in an ophthalmic drug composition. It is used, together with insulin like growth factor-I (IGF-I), to treat corneal disorders such as corneal ulcer, corneal epithelial peeling, dry eye and Keratitis. The dosage is 0.1-5000 (preferably 1-1000) mg/day of the tetrapeptide and 0.001-100 (preferably 0.01-10) mg/day of IGF-I. The preferable form of the composition is eye drops

Sequence 4 AA;

Human/rat tachykinin Substance P C-terminal tetrapeptide.

16-JAN-2001 (first entry)

AAB23026;

AAB23026 standard; peptide; 4 AA.

AAB23026 RESULT

FGLM

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This invention describes a novel aqueous suspension of monodisperse particles on non-crosslinked, non-denatured albumin (50-5000 nm) which is stable against dissolving upon dilution with an alcohol-free aqueous candium. The method involves (a) forming an aqueous solution containing albumin and hemoglobin and (b) treating the aqueous solution containing a alcohol to cause the solution to become turbid. The particles are useful as agents for in vivo administration, either of their own administration or as a vehicle for other therapeutic or diagnostic agents. The method nameter and micrometer size range, in a form closer to their natural form than the forms of the prior art. The particles in the form of a more closely controlled agent for in vivo administration, with greater a more closely controlled agent for in vivo administration, with greater case of clearance from the body after their period of usefulness.
                                             ö
                                          Gaps
                                                                                                                                                                                                                                                                         Non-crosslinked protein particle; diagnostic; therapy; monodisperse; albumin; haemoglobin; nanometer; micrometer; clearance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Non-crosslinked protein particles for therapeutic and diagnostic use,
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0
            Length 4;
                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 21; DB 2; Length 4; Conservative 0; Mismatches 0; Indels
    100.0%; Score 21; DB 2; L
100.0%; Pred. No. 1.8e+06;
Live 0; Mismatches 0;
                                                                                                                                                                                                                                             Non-crosslinked protein particle peptide 124.
                                                                                                                                                                                                                                                                                                                                                                               /note= "C-terminal amide"
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                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                          AAY31075 standard; peptide; 4 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          91US-00641720.
92US-00959560.
93US-00069831.
94US-00212546.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 96US-00747137
                                                                                                                                                                                                               21-OCT-1999 (first entry)
                                4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (HEMO-) HEMOSPHERE INC.
Query Match
Best Local Similarity
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Modified-site
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01-JUN-1993;
14-MAR-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                      USS945033-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-JAN-1991;
                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                        AAY31075;
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                              Matches
                                                                                                                                RESULT 2
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The inevntion relates to a method for detecting magnesium binding defects. The method comprises quantitating a tachykinin C-terminal Phatapeptide (e.g., AMB23025) and its degradation products (e.g., AMB23026) in blood using an antibody specific for the generalises mammalian tachykinin C-terminal pentapeptide Phe-(Phe/Val)-Gly-Leu-Met-MHZ (AAB23028). The method is useful for detecting callular magnesium binding defects which are associated with abnormal physiological states such as sodium-sensitive essential hypertension and type 2 diabetes mellitus. The present sequence represents the C-terminal 4 amino acids of the tachykinin Substance P (AAB23027) from human and rat. This is a degradation product of the Substance P C-terminal pentapeptide (AAB23025) and may also be assayed according to the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detecting magnesium binding defects associated with abnormal physiological states such as sodium-sensitive essential hypertension and type 2 insulin-resistant diabetes mellitus, comprises measuring a specific pentapeptide in blood.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                          Substance P; tachykinin; human; rat; magnesium binding defect; sodium sensitive essential hypertension; inmulin resistance; type 2 diabetes; antibody; immunoassay; quantification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ó
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "C-terminal amide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 5; 21pp; English.
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Modified-site
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Thibaudeau K;

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The present invention describes a modified therapeutic peptide (I) comprising a therapeutically active amino acid region (III) and a reactive group (II) (e.g. succinimidyl and maleimido groups) attached to a less therapeutically active amino acid region (IV), which covalently bonds with amino/hydroxyl/thiol groups on blood components to form a peptidase stabilised therapeutic peptides composed of 3-50 amino acids. (I) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity in vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body.

Modifying and attaching therapeutic peptides to albumin prevents or reduces the action of peptidases to increase length of activity (half iffe) and specificity as bonding to large molecules decreases.

Albeboas to Arbeztal represent peptides which can be used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wound healing; insulin-like growth factor-I; tear; abrasion; skin ulcer; surgical incision; burn.
                                                                                                                                                                                                                                                                                                    Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 21; DB 4; Length 4; 100.0%; Pred. No. 1.8e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Substance P analog used in wound healing treatment#14.
                                                                                                                                                                                                                    Bridon DP, Ezrin AM, Milner PG, Holmes DL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 402; 733pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABB10091 standard; peptide; 4 AA
                                                                                              99US-0134406P.
99US-0153406P.
99US-0159783P.
                                                        L7-MAY-2000; 2000WO-US013576.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-AUG-2001; 2001WO-JP006933.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-AUG-2000; 2000JP-00242489.
28-NOV-2000; 2000JP-00361388.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4; Conservative
                                                                                                                                                                              (CONJ-) CONJUCHEM INC.
                                                                                                                                                                                                                                                          WPI; 2001-112059/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 FGLM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Inidentified.
                                                                                                17-MAY-1999;
                                                                                                                 10-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-FEB-2002.
                  23-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABB10091.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention provides a pharmaceutical and/or veterinary formulation that comprises 2 -30% of active agents which include a gonadorropin-releasing hormone (GnRH) agonist, 0.5 - 20% of a pore-forming agent which is not lecithin, and the remainder stearin. The formulation is useful as a sustained release implant which can deliver the active agent for a period of 7 days to 2 years. Sequences AAV67573-578 represent P antagonist peptides used in the composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bioimplant formulation for sustained delivery of an active agent over 7 days to 2 years, comprises active agent, pore-forming agent and stearin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Protection; endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter.
                                                                                                                   Pharmaceutical, veterinary, gonadotropin-releasing hormone, GnRH, pore-forming agent; lecithin, stearin, P antagonist.
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                                                                                                                                                                                                                                                            /note= "C-terminal amide"
                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fachykinins peptide SEQ ID NO:623.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rathjen DA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 20; Page 21; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB91447 standard; peptide; 4 AA.
                                                                                                                                                                                                                                                                                                                                                                                    99WO-AU000585.
                                                                                                                                                                                                                                                                                                                                                                                                                            98AU-00004730.
                                                                                                                                                                                                                                                                                                                                                                                                                                             98AU-00004731
                                       19-MAY-2000 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4; Conservative
                                                                             P antagonist peptide #5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Trigg TE, Walsh JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-182528/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (PEPT-) PEPTECH LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 FGLM 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 4 AA;
                                                                                                                                                                                                                                                                                                    WO200004897-A1
                                                                                                                                                                                                                                                                                                                                                                                  20-JUL-1999;
                                                                                                                                                                                                                        Key
Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                             20-JUL-1998;
13-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                          20-JUL-1998;
                                                                                                                                                                              Unidentified
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Synthetic.
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AAB91447;

RESULT 5 AAB91447

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Gaps

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The invention relates to skin wound healing promoters, containing substance P analogs or their pharmaceutically-acceptable salts, and insulin-like growth factor-I as the active ingredient. The promoters are for treating wounds like tears, abrasions, surgical incisions, or skin ulcers and burns. The current sequence represents a substance P analog for use in wound healing treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tachykinin; substance P; hypertension; hypotensive; antidiabetic; gynaecological; salt-insensitive hypertension; magnesium binding; insulin resistance; type 2 diabetes mellitus; pre-eclampsia; eclampsia.
                                                                                        Skin wound healing promoters or skin epidermal extension promoters containing substance P analogs and insulin-like growth factor-I for treating wounds like tear, abrasion, surgical incision, skin ulcers burns.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New monopoptides derived from butadienes, ethylenes and propanes are magnesium binding defect antagonists, useful in the treatment of e.g. hypertension, insulin resistance of type 2 diabetes mellitus and
                                                                                                                                                                                                                                                                                                        100.0%; Score 21; DB 5; Length 4; 100.0%; Pred. No. 1.8e+06; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4. .4
/note= "C terminal-amide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tachykinin N -terminal tetrapeptide.
                                           Nakamura M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                        Claim 3; Page 11; 20pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MAGN-) MAGNESIUM DIAGNOSTICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                AAU77846 standard; peptide; 4 AA.
(SANT ) SANTEN PHARM CO LTD. (NISH/) NISHIDA T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-AUG-2000; 2000US-00635266.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-AUG-2001; 2001WO-US024909
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                 4; Conservative
                                    Nishida T, Nakata K,
                                                                WPI; 2002-269153/31.
                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-280663/32.
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                                                                                                                                                                                                                                                                              Sequence 4 AA;
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                                                                                                                                                                                                                                                                                                       Query Match
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Matches
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Disclosure; Page 2; 38pp; English,

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Gaps

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This invention relates to novel therapeutic compounds and methods used
for for treating mammals with disorders such as salt-insensitive
hypertension. The monopepited compounds of the invention are derived from
bladdenss, ethylenes and propanes. The compounds of the invention are
concentration of magnesium binding within the plasma membranes
of somatic cells which results in a decrease in the intracellular
concentration of magnesium ions. These compounds may be used in the
treatment of a mammal affected with magnesium binding defect, salt-
consistive (particularly hypertension), insulin resistance of type 2
diabetes mellitus and pre-eclampsia/cclampsia The compounds of the
compounds are biologically stable. The present sequence represents the a
compounds are biologically stable. The present sequence represents the a
compounds are biologically stable. The present sequence represents the a
cubstance P, this peptide is sufficient to orrect the magnesium binding
                                                                                                                                                                                                                                                                                                                                                                                ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel peptides based on minimum activity expression units of insulin-like growth factor-1, applicable in remedies in ophthalmology and dermatology for treating keratic injury and promoting wound healing in skin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to the determination of the smallest peptide fragment of insulin growth factor 1 (IGF-1) with the highest activity for use in ophthalmology and dermatology. The peptides are applicable in remedies in ophthalmology and dermatology for treating keratic injury and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ophthalmological; dermatological; vulnerary; insulin growth factor 1; IGF-1; ophthalmology; dermatology; keratic injury; wound healing; skin; corneal ulcer; exofoliation of corneal epithelium; keratitis; dry eye; scratch; surgical cutting; skin ulcer; burns.
                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               High activity minimal IGF-1-derived peptide fragment #10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "amidated C-terminus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADE94198 standard; peptide; 4 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nakamura M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-DEC-2002; 2002WO-JP012632
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nishida T, Inui M,
                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                            Sequence 4 AA;
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Assessing predisposition to physiological disorder associated with magnesium binding defect in individual, by measuring level of amidated peptides associated with magnesium binding defect in sample and comparing
promoting wound healing in the skin. The keratic injury is particularly corneal ulcer, exofoliation of corneal epithelium, keratitis or dry eye. The skin wound can be scratches, surgical cutting, skin ulcer, or burns. This sequence represents one of the peptides of the invention with IGF-1
                                                                                                                                                                                                                                                                                                                                                                            physiological disorder; preeclampsia; pregnancy;
salt-sensitive essential hypertension; type 2 diabetes mellitus; human.
                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                             Magnesium binding defect; MgBD; MgBD binding defect peptide mimetic;
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                                                                                                       100.0%; Score 21; DB 7; Length 4; 100.0%; Pred. No. 1.8e+06;
                                                                                                                                                                                                                                                                                                                                   Human magnesium binding defect (MgBD) peptide mimetic #2.
                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "OTHER= C-terminal amide"
                                                                                                                                0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                     ADR43772 standard; peptide; 4 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /label= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-AUG-2000; 2000US-00635266.
24-JAN-2002; 2002US-00053669.
29-AUG-2002; 2002US-00230133.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-MAR-2004; 2004US-00805881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99US-00265690
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-OCT-2003; 2003US-00695536
                                                                                                                                                                                                                                                                                                          18-NOV-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           peptide level to standard.
                                                                                                                                   4; Conservative
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                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (WELL/) WELLS I C.
                                                                                                                                                            1 FGLM 4
                                                                                                                                                                                   FGLM 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US2004171093-A1
                                                                              Sequence 4 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                                                                                                                                                                                                                                                                 ADR43772;
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                                                    activity
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peptide after treatment, where a significant increase in the level of the progress of treatment of the individual, a monoclonal antibody that specifically binds to a peptide or its peptide minetic, a prognosis reagent for determining the presence of a magnesium conclonal antibody that specifically binds to a peptide or its peptide or minetic, a prognosis reagent for determining the presence of a magnesium conclonal mammalian somatic cells involving obbaining a sample of magnesium in mammalian somatic cells, collecting the somatic cells from the body fluid by centrifigation, resuspending the somatic cells from the body fluid by centrifigation, resuspending the somatic cells from the body fluid by centrifigation, resuspending the somatic cells and repeating the removing and mesuring steps at collectify measuring the level of tightly bound magnesium is somatic cells and repeating the removing and mesuring steps at significantly reduced and the somatic cells remain intact, a method of identifying substances which promote binding of tightly bound magnesium in a physiological medium including magnesium in a physiological medium including magnesium in a physiological medium including magnesium ion, adding a substance to be tested to the suspension and measuring the level of the substance to be tested to the suspension and measuring the level of bound magnesium after addition of the substance to be tested is ciphtly bound magnesium after addition of the substance to be tested is involving administering to the individual a substance which promotes to involving administering to the individual a substance which promotes to predisposition to a physiological disorder associated with a magnesium control of physical disorder associated with the magnesium control of physical disorder associated with the magnesium control of the invention. This sequence represents a human was mentioned is also useful for ameliorating or correcting a magnesium or correcting a magnesium or correcting a magnesium or correcting a magnesium and an a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 21; DB 8; Length 4; 100.0%; Pred. No. 1.8e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cell growth inhibitor; tumour cell growth inhibitor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /label= Carbobenzoxy-Phe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAP61654 standard; peptide; 4 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              83US-00552255.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
(revised)
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 FGLM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-NOV-1983;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAP61654;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Gaps

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This sequence is shown in the specification. The invention relates to an ophthalmic drug composition which contains Phe-Gly-Leu-Met-NH2 or its medicinally acceptable salts as the active ingredient. It is used, together with innulin like growth factor—I (IGF-I), to treat corneal disorders such as corneal ulcer, corneal epithelial peeling, dry eye and keratitis. The dosage is 0.1-5000 (preferably 1-1000) mg/day of the active ingredient and 0.001-100 (preferably 1-1000) mg/day of the preferable form of the composition is eye drops
                                                                                                                                                                                                                                                                                                                         Ophthalmic drug; corneal disorder; ulcer; epithelial peeling; dry eye;
keratitis; insulin like growth factor-I; IGF-I; eye drop.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ophthalmic drug composition containing tetra:peptide - is useful as corneal disorder remedy for corneal ulcer, corneal epithelial peeling, dry eye, keratitis.
        Score 18; DB 1; Length 4;
Pred. No. 1.8e+06;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     85.7%; Score 18; DB 2; Length 4;
75.0%; Pred. No. 1.88+06;
.ive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "C-terminal amide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 11; 19pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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                                                                                                                                                                                      AAW41686 standard; peptide; 4 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nakata
       85.7%;
ilarity 75.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 96JP-00165612.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SANT ) SANTEN PHARM CO LTD.
                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3; Conservative
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 Query Match
Best Local Similarity
Matches 3; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1998-076907/07.
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Matches 3, Conserv
                                                                         4
                                                                                                       4
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                                                                       FGLM
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Modified-site
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1 YGLM 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 4 AA;
                                                                                                                                                                                                                                                        09-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9749419-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-DEC-1997.
                                                                                                                                                                                                                        AAW41686;
                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                      RESULT 12
                                                                                                                                                                        AAW41686
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ID ABB1
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AC ABB1
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                                                                                                                                                                                                                                 The inventors
                                                                                                                                                                                                                 The cpds. of the invention inhibit protein phosphorylation. The inventors claim a process for inhibiting the growth of tumour cells in a medium which comprises contacting the cells with a cpd. of formula (AAP61654-P61661) or a physiologically acceptable salt. (Updated on 03-0CT-2002 to add missing OS field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This peptide is useful as part of a peptide/halo-methyl ketone cpd., for treating picornavirus, egpolio- or rhinovirus infections. It inhibits the processing of picornavirus capsid proteins by virus encoded proteases. See AAP71302-13. See also US4652552. (Updated on 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Treating picorna-virus infection with peptide halo.methyl ketone cpds.
esp. for treating polio virus and rhino virus infections.
                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                  Inhibition of tumour cell growth - using peptide and aminoacid halo:methyl ketone(s).
                                                                                                                                                                                                                                                                                                                                                                                                     ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptide component of cpd. for treating picornavirus infections.
                                                                                                                                                                                                                                                                                                                                                       / Match 85.7%; Score 18; DB 1; Length 4; Local Similarity 75.0%; Pred. No. 1.8e+06; nes 3; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Picornaviridae, poliovirus, rhinovirus, antiviral agent.
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                              (DUPO ) DU PONT DE NEMOURS & CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAP71301 standard; peptide; 4 AA.
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   83US-00552255.
                                                                                                                                                                                  Claim 1; Col 4; 9pp; English.
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (DUPO ) DU PONT DE NEMOURS
                                                                  Kettner CA, Racker E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kettner CA, Korant BD;
                                                                                                  WPI; 1986-118872/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1987-036897/05.
                                                                                                                                                                                                                                                                                                                                                                                                                           1 FGLM 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                |||:
1 FGLL 4
                                                                                                                                                                                                                                                                                                                         Sequence 4 AA;
16-NOV-1983;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAR-2003
15-MAY-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
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Query Match Best Loca Matches

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RESULT 11

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Gaps

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ABB10092

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The cpds. of the invention are useful in diagnosis of infectious diseases caused by viruses which encode a specific protease e.g. picornaviruses. (Updated on 0-OCT-2002 to add missing OS field.) (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Treating picorna-virus infection with peptide halo:methyl ketone cpds. esp. for treating polio virus and rhino virus infections.
             /note= "Bonded to a chromogenic, fluorogenic, chemiluminescent, radioactive, antigenic, or haptenic indicator group."
                                                                                                                                                                                                                                                                                                                    Peptide substrates for virus-specified protease(s) - with C-terminal indicator gp. linked by amide or ester linkage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peptide component of cpd. for treating picornavirus infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        76.2%; Score 16; DB 1; Length 4; 100.0%; Pred. No. 1.8e+06; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Picornaviridae; poliovirus; rhinovirus; antiviral agent
                                                                                                                                                                                                                   (DUPO ) DU PONT DE NEMOURS & CO E I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (DUPO ) DU PONT DE NEMOURS & CO E I.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAP71312 standard; protein; 4 AA.
                                                                                                                                                  86EP-00300147.
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                                                                                                                                                                                                                                                                                                                                                                     Example; p22; 41pp; English.
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Best Local Similarity 100.00
Thes 3; Conservative
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(first entry)
                                                                                                                                                                                                                                                    Korant BD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1987-036897/05.
                                                                                                                                                                                                                                                                                   WPI; 1986-184617/29
 Misc-difference
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                                                                                                                                                  10-JAN-1986;
                                                                                                                                                                                   11-JAN-1985;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-AUG-1984;
                                                                                                                                                                                                                                                    Kettner CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-MAR-2003
15-MAY-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US4636492-A
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                                                                                  EP187721-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAP71312;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "bonede to Boc, Z, Suc, or MeOSuc; Z=carbobenzoxy;
Boc=t-Butyloxycarbonyl; Suc=Succinyl;
MeOSuc=Methoxysuccinyl"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to skin wound healing promoters, containing substance P analogs or their pharmaceutically-acceptable salts, and insulin-like growth factor-I as the active ingredient. The promoters are for treating wounds like tears, abrasions, surgical incisions, or skin ulcers and burns. The current sequence represents a substance P analog for use in wound healing treatment
                                                                         Wound healing; insulin-like growth factor-1; tear; abrasion; skin ulcer; surgical incision; burn.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Skin wound healing promoters or skin epidermal extension promoters containing substance P analogs and insulin-like growth factor-I for treating wounds like tear, abrasion, surgical incision, skin ulcers or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence located immediately adjacent to and upstream of the cleavage site within a virus-specified polypeptide precursor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 4;
                                               Substance P analog used in wound healing treatment#15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 18; DB 5; I
Pred. No. 1.8e+06;
1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Viral desease; diagnosis; picornavirus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 4; 20pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                       Nakamura M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAP61707 standard; peptide; 4 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          85.7%; initiating 15.0%; in Conservative 1,1
                                                                                                                                                                                                                                                                10-AUG-2000; 2000JP-00242489.
28-NOV-2000; 2000JP-00361388.
                                                                                                                                                                                                                                  10-AUG-2001; 2001WO-JP006933
                                                                                                                                                                                                                                                                                                                   (SANT ) SANTEN PHARM CO LTD. (NISH/) NISHIDA T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
               26-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
                                                                                                                                                                                                                                                                                                                                                                     Nishida T, Nakata K,
                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-269153/31
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1 YGLM 4
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Misc-difference
                                                                                                                                                                WO200213853-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 4 AA;
                                                                                                                                 Unidentified.
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03-OCT-2002
08-JUN-1991
                                                                                                                                                                                               21-FEB-2002
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Synthetic

AAP61707;

RESULT 14

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Query Match Best Loc Matches

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Gaps

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X222222X

This peptide is useful as part of a peptide/halo-methyl ketone cpd., for treating picornavirus, egpolio- or rhinovirus infections. It inhibits the processing of picornavirus capsid proteins by virus encoded proteases. See AAP71301-11 and AAP71313. See also US4652552. (Updated on 25-MAR-2003 to correct PA field.) Sequence 4 AA;

Query Match 76.2%; Score 16; DB 1; Length 4; Best Local Similarity 100.0%; Pred. No. 1.8e+06; Matches 3; Conservative 0; Mismatches 0; Indels ò

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Search completed: March 23, 2005, 15:12:55 Job time : 167 secs

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March 23, 2005, 15:13:04; Search time 137 Seconds Run on:

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BLOSUM62 Scoring table:

Gapop 10.0 , Gapext 0.5

1407402 seqs, 331100923 residues Searched:

9312 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 4

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Published Applications AA:* Database :

| Cgn2_6/ptodata1/lpubpaa/US07_PUBCOMB.ppp:*
| Cgn2_6/ptodata1/lpubpaa/US07_PUBCOMB.ppp:*
| Cgn2_6/ptodata1/lpubpaa/US07_NEW_PUB.ppp:*
| Cgn2_6/ptodata1/lpubpaa/US06_PUBCOMB.ppp:*
| Cgn2_6/ptodata1/lpubpaa/US06_PUBCOMB.ppp:*
| Cgn2_6/ptodata1/lpubpaa/US07_NEW_PUB.ppp:*
| Cgn2_6/ptodata1/lpubpaa/US08_PUBCOMB.ppp:*
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| Cgn2_6/ptodata1/lpubpaa/US09_NEW_PUB.ppp:*
| Cgn2_6/ptodata1/lpubpaa/US09_NEW_PUB.ppp:*
| Cgn2_6/ptodata1/lpubpaa/US10_PUBCOMB.ppp:*
| Cgn2_6/ptodata1/lpubpaa/US0_NEW_PUB.ppp:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Description	Sequence 2, Appli	Sequence 3, Appli	Sequence 2, Appli	Seguence 3, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 9, Appli	Seguence 270, App	Sequence 2, Appli	Sequence 2, Appli	Sequence 8, Appli	Sequence 328, App	Sequence 98, Appl
ΩI	US-09-265-690C-2	US-10-230-133-3	US-10-053-669-2	US-10-695-536-3	US-10-805-881-2	US-10-497-628-2	US-09-879-442A-9	US-10-821-240A-270	US-10-230-133-2	US-10-695-536-2	US-09-879-442A-8	US-10-137-867-328	US-09-879-442A-98
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Query Match	100.0	100.0	100.0	100.0	100.0	100.0	76.2	76.2	71.4	71.4	66.7	66.7	61.9
Score	21	21	21	21	21	21	16	16	15	15	14	14	13
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Sequence 99, Appl Sequence 24, Appl Sequence 30, Appl Sequence 30, Appl	2224 201111989 2024, 4, 4, 4, 4, 4, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,	Sequence 13, Appl
4 9 US-09-879-442A-99 4 9 US-09-943-123-24 4 14 US-10-087-905-30 4 14 US-10-087-942-30 4 14 US-10-087-402-30	10.108-3894 10.202-824 10.202-824 10.359-393 10.121-857 10.121-857 10.255-679 10.255-679 10.007-884-409- 08-884-409- 08-884-409- 09-811-945 10.007-6811-945 10.007-007-611-007-005 10.007-005-679 10.007-005-679	4 14 US-10-255-679-13
14 13 61.9 15 13 61.9 16 13 61.9 17 13 61.9	23 23 23 23 24 25 25 25 25 25 25 25 25 25 25 25 25 25	45 12 57.1

US-09-265-690C-2
US-09-265-690C-2
Sequence 2, Application US/09265690C
Sequence 2, Application US/09265690C
Sequence 2, Application No. US20010051345A1
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Method for Detecting Deficient Cellular Membrane Tightly Bound Mac
TITLE OF INVENTION: for Disease Diagnosis
TITLE OF INVENTION: for Disease Diagnosis
TITLE OF INVENTION: 102001
CURRENT APPLICATION NUMBER: US/09/265,690C
CURRENT FILING DATE: 1999-03-10
SOFTWARE: PatentIn version 3.0
SOFTWARE: PatentIn version 3.0 LENGTH:

ALIGNMENTS

TYPE: PRT

ORGANISM: Homo sapiens

FBATURE:

AMME/KEY: MOD_RES

LOCATION: (4) ... (4)

OTHER INFORMATION: AMIDATION
US-09-265-690C-2

Gaps ; 100.0%; Score 21; DB 9; Length 4; 100.0%; Pred. No. 1.3e+06; tive 0; Mismatches 0; Indels Query Match Best Local Similarity 100. Matches 4; Conservative

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1 FGLM 4 ઠે g RESULT 2 US-10-230-133-3 Sequence 3, Application US/10230133 ; Publication No. US20030040625A1

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US-10-805-801-2

Sequence 2, Application US/10805881

Sequence 2, Application Work US-20040171093A1

Sequence 2, Application NO. US20040171093A1

SEQUENCEANT: Wells, IDext C.

TITLE OF INVENTION: Method for Detecting Deficient Cellular Membrane Tightly Bound

TITLE OF INVENTION: Magnesium for Disease Diagnosis

FILE REFERENCE: 800812-0005

CURRENT FILING DATE: 2004-03-22

PRIOR APPLICATION NUMBER: US/10/695,881

PRIOR APPLICATION NUMBER: US 10/653,669

PRIOR APPLICATION NUMBER: US 10/655,536

NUMBER OF SEQ ID NOS: 4

SEQ ID NO 2

SEQ ID NO 2

LENGTH: 4
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APPLICANT: Masatsugu Nakamura
TITLE OF INVENTION: NOVEL PEPTIDE AND PHARMACEUTICAL USE OF THE SAME
FILE REFERENCE: 04355/HG
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100.0%; Pred. No. 1.3e+06;
iive 0; Mismatches 0;
                                                              CURRENT FILING DATE: 2003-10-28
PRIOR APPLICATION NUMBER: US 10/230,133
PRIOR FILING DATE: 2002-08-29
PRIOR FILING DATE: 2002-08-29
PRIOR FILING DATE: 2002-08-29
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin version 3.2
LENGTH: 4
                                             CURRENT APPLICATION NUMBER: US/10/695,536
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Publication No. US20050009752A1
GENERAL INFORMATION:
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FEATURE:
NAMMEKEY: MOD RES
LOCATION: (4) ... (4)
OTHER INFORMATION: AMIDATION
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; OTHER INFORMATION: AMIDATION
US-10-805-881-2
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 4; Conserv
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                                                       TITLE OF INVENTION: Antagonists of the magnesium binding defect as therapy agents and TITLE OF INVENTION: Antagonists of the magnesium binding defect as therapy agents and TITLE OF INVENTION: methods for treatment of abnormal physiological states FILE OF INVENTION: methods for treatment of abnormal physiological states CURRENT PAPLICATION NUMBER: US/10/230,133 PRIOR PRILING DATE: 2002-08-29 PRIOR FILING DATE: 2000-08-09 PRIOR FILING DATE: 2000-08-09 PRIOR FILING DATE: 2000-08-09 PRIOR FILING DATE: 2000-08-09 PRIOR PRILING DATE: 2000-08-09 PRIOR PRINCE PRINCE
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TITLE OF INVENTION: Method for Detecting Deficient Cellular Membrane Tightly Bound Ma
TITLE OF INVENTION: for Disease Diagnosis
FILE REFERENCE: N1427-005
GURRENT APPLICATION NUMBER: US/10/053,669
FRIOR APPLICATION NUMBER: 09/265,690
PRIOR APPLICATION NUMBER: 09/265,690
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin version 3.0
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TITLE OF INVENTION: Antagonists of the Magnesium Binding Defect as Therapeutic Agents
TITLE OF INVENTION: and Methods for Treatment of Abnormal Physiological States
FILE REFERENCE: 800812-0008
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100.0%; Score 21; DB 14; Length 4;
Best Local Similarity 100.0%; Pred. No. 1.3e+06;
Matches 4; Conservative 0; Mismatches 0; Indels
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100.0%; Pred. No. 1.3e+06;
iive 0; Mismatches 0;
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Publication No. US20030077658A1
GENERAL INFORMATION:
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Sequence 3, Application US/10695536
Publication No. US20040110692A1
GENERAL INFORMATION:
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; LOCATION: (4)...(4)
; OTHER INFORMATION: AMIDATION
US-10-230-133-3
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; LOCATION: (4)...(4)
; OTHER INFORMATION: AMIDATION
US-10-053-669-2
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Best Local Similarity 100.
Matches 4; Conservative
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
GENERAL INFORMATION:
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OTHER INFORMATION: Description of Artificial Sequence: Synthetic
NAME/KEY: SITE
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APPLICANT: Lewis, Evan
APPLICANT: Nieder, Matthew H.
APPLICANT: Pickford, Lesley B.
APPLICANT: Pickford, Lesley B.
APPLICANT: Trouet, Andre
APPLICANT: Yarranton, Geoffrey T.
TITLE OF INVENTION: ENZYME CLEAVABLE PRODRUG COMPOUNDS
FILE. REPERENCE: COUL-015/0208
CURRENT APPLICATION NUMBER: US/09/879,442A
                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR PELICATION NUMBER: 60/290,448
PRIOR FILING DATE: 2001-05-11
PRIOR APPLICATION NUMBER: 60/211,887
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 1999-12-10
PRIOR PELING DATE: 1999-02-08
PRIOR PELING DATE: 1999-02-08
PRIOR PELING DATE: 1999-12-10
PRIOR PELING DATE: 1999-12-10
PRIOR PELING DATE: 1998-12-11
PRIOR PELING DATE: 1998-12-11
PRIOR FILING DATE: 1998-12-11
PRIOR FILING DATE: 1998-12-11
PRIOR FILING DATE: 1998-12-11
CURRENT APPLICATION NUMBER: US/10/497,628
CURRENT FILING DATE: 2004-06-03
PRIOR APPLICATION NUMBER: JP 2001-368103
PRIOR FILING DATE: 2001-12-01
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PATENTLY VET: 2.1
SEQ ID NO 2
LENGTH: 4
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Fernandez, Anne Marie
Gangwar, Sanjeev
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; LOCATION: (1)
; OTHER INFORMATION: Beta-Alanine
US-09-879-442A-9
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ORGANISM: Artificial Sequence
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Best Local Similarity
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Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                                                              1 FGLM 4
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                                                                                                                                                                                 ; TYPE: PRT
; ORGANISM: Human
US-10-497-628-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -09-879-442A-9
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RESULT 8

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FEATURE:
) OTHER INFORMATION: Description of Artificial Sequence: derivative peptide based on mail to the Court of the Cou
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TITLE OF INVENTION: Antagonists of the magnesium binding defect as therapy agents and
TITLE OF INVENTION: Antagonists of treatment of abnormal physiological states
FILE REFERENCE: 2892-106
CURRENT APPLICATION NUMBER: US/10/230,133
PURRENT PILING DATE: 2002-08-29
PRIOR APPLICATION NUMBER: 09/635,266
PRIOR FILING DATE: 2000-08-09
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SQUENCE 270, Application US/10821240A
Publication No. US20050037430A1
GENERAL INPORMATION:
GENERAL INPORMATION:
TITLE OF INVENTION: Gene regulator
TITLE OF INVENTION: Gene regulator
FILE REFERENCE: 2183-5223U8
CURRENT APPLICATION NUMBER: US/10/821,240A
CURRENT FILING DATE: 2004-04-08
PRIOR FILING DATE: 2001.12-21
PRIOR FILING DATE: 2001.12-21
PRIOR FILING DATE: 2001.10-04
NUMBER OF SEQ ID NOS: 312
SEQ ID NO 270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
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Publication No. US20030040625A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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SOFTWARE: Patentin version 3.0
SEQ ID NO 2
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NAME/KEY: MOD RES
LOCATION: (3) ... (3)
COTHER INFORMATION: AMIDATION
US-10-230-133-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.
Matches 3; Conservative
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Best Local Similarity 100.
Matches 3; Conservative
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ORGANISM: Homo sapiens
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Sequence 328, Application US/10137867
Publication No. US20030207349A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dubois, Vincent
Fernandez, Anne Marie
Gangwar, Sanjeev
Lewis, Evan
                 66.7%;
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Gerritsen, Mary E.
Goddard, Audrey
                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Beresini, Maureen
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            Query Match
Best Local Similarity 66.7
Matches 2; Conservative
                                                                                                                                                                                                                                                                                                              Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
                                                                                                                                                                                                                                             Desnoyers, Luc
Filvaroff, Ellen
                                                                                                                                                                                                                                                                                                                                                        Smith, Victoria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2; Conservative
                                                                                                                                                                                                                                   DeForge, Laura
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ORGANISM: Homo Sapien
US-10-137-867-328
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Best Local Similarity
Matches 2; Conserv
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       TITLE OF INVENTION: Antagonists of the Magnesium Binding Defect as Therapeutic Agents TITLE OF INVENTION: Antagonists of the Magnesium Binding Defect as Therapeutic Agents FILE OF INVENTION: and Methods for Treatment of Abnormal Physiological States CURRENT APPLICATION NUMBER: US/10/695,536

CURRENT APPLICATION NUMBER: US/230,133

PRIOR APPLICATION NUMBER: US 09/635,266

PRIOR APPLICATION NUMBER: US 09/635,266

NUMBER OF SEQ ID NOS: 4

SOFTWARE: Patentin version 3.2

SEQ ID NO 2

SEQ ID NO 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 3;
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  Clifton
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                                                                                                                                                                                                                           ; NAME/KEY: MOD_RES
; LOCATION: (3)...(3)
; OTHER INFORMATION: AMIDATION
US-10-695-536-2
                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.
Matches 3; Conservative
APPLICANT: Wells, Ibert
                                                                                                                                                                                     TYPE: PRT
ORGANISM: Homo sapiens
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CURRENT APPLICATION NUMBER: 2002-05-03

Prior Application removed - See Palm or File Wrapper NUMBER OF SEQ ID NOS: 550

SEQ ID NO 328

LENGTH: 379

TYPE: ....
                                                            Gaps
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0
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Score 14; DB 9; Length 4;
Pred. No. 1.3e+06;
1; Mismatches 0; Indels
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APPLICANT: Lobl, Thomas J.
APPLICANT: Lobl, Thomas J.
APPLICANT: Pickford, Lesley B.
APPLICANT: Prouet, Andre
APPLICANT: Trouet, Andre
APPLICANT: Tranet, Andre
APPLICANT: Tranet, Geoffrey T.
TITLE OF INVENTION: ENZYME CLEAVABLE PRODRUG COMPOUNDS
FILE REPREBENCE: COUL-015/02US
CURRENT APPLICATION NUMBER: US/09/879,442A
CURRENT FILING DATE: 2001-06-11
PRIOR APPLICATION NUMBER: 60/290,448
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NAME/KEY: SITE
LOCATION: (1)
COTHER INFORMATION: 2-Thienylalanine
US-09-879-442A-98
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APPLICANT: Trouget, Andre
APPLICANT: Trouget, Andre
TILEO TINVENTION, GGOFFREY T.
FILE REFERENCE: COUL-015/0203
CURRENT APPLICATION NUMBER: US/09/879,442A
CURRENT FILING DATE: 2001-06-11
PRIOR FILING DATE: 2001-06-11
PRIOR FILING DATE: 2001-06-14
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 1999-12-10
PRIOR APPLICATION NUMBER: 60/119,312
PRIOR APPLICATION NUMBER: 60/119,312
PRIOR APPLICATION NUMBER: 60/119,312
PRIOR PILING DATE: 1999-02-08
PRIOR APPLICATION NUMBER: 60/119,312
PRIOR FILING DATE: 1999-12-11
NUMBER OF SEQ ID NOS: 103
SOFTWARE: PATENTIN OFF: 2.1
PRIOR FILING DATE: 2001-05-11
PRIOR APPLICATION NUMBER: 60/211,887
PRIOR PILING DATE: 2000-06-14
PRIOR FILING DATE: 1000-06-14
PRIOR FILING DATE: 1999-12-10
PRIOR PILING DATE: 1999-12-10
PRIOR APPLICATION NUMBER: 60/119,312
PRIOR FILING DATE: 1999-02-08
PRIOR FILING DATE: 1998-12-11
NUMBER OF SEQ ID NOS: 103
SOFTWARE: PATENTIN VEY: 2.1
SEQ ID NO 98
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US-09-879-442A-99
; Sequence 99, Application US/09879442A
; Patent US200142955A1
; GENERAL INFORMATION:
; APPLICANT: CORIXA CORPORATION
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Fernandez, Anne Marie
Gangwar, Sanjeev
Lewis, Evan
Lobl, Thomas J.
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Pickford, Lesley B.
                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
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2 YGL 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 FGL 3
                                                                                                                                                                                                                                                                                                         TYPE: PRT
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APPLICANT: CHANG, Y-H
APPLICANT: VETRO, J.A.
APPLICANT: WETRO, J.A.
APPLICANT: MICKA, W.S.
TITLE OF INVENTION: 2 ("MetAP2") and Clinical Uses Therefor
TITLE OF INVENTION: 2 ("MetAP2") and Clinical Uses Therefor
FILE REPERENCE: 16153-8007
CURRENT APPLICATION NUMBER: US/09/943,123
CURRENT FILING DATE: 2001-08-30
NUMBER: OF SEQ ID NOS: 26
SOFTWARE: PatentIn Ver: 2.0
SEQ ID NO 24
LENGTH: 4
                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: synthetic
OTHER INFORMATION: peptide
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Score 13; DB 9; Length 4;
Pred. No. 1.3e+06;
1; Mismatches 0; Indels
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                                                                                                                                                                                         RESULT 15
US-09-943-123-24
Sequence 24, Application US/09943123
Publication No. US20020182701A1
  61.9%;
  Query Match
Best Local Similarity 66.7
Matches 2; Conservative
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Best Local Similarity 66.7
Matches 2; Conservative
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                                                                                        1 FGL 3
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US-08-441-591-63
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Sequence 20, Appli
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Sequence 4, Appli
Sequence 25, Appli
Sequence 25, Appli
Sequence 60, Appli
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Sequence 63, Appl
Sequence 2, Appli
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Sequence 3, Appli
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                                                                                                                    March 23, 2005, 15:04:03; Search time 41 Seconds (without alignments)
7.283 Million cell updates/sec
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Sequence 31
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/cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PcTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PcTUS_COMB.pep:*
GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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US-08-303-362A-63
US-09-265-607-2
US-09-265-607-2
US-10-230-133-3
US-10-230-133-3
US-08-747-137-124
US-08-722-126A-20
US-09-635-266-2
US-09-635-266-2
US-08-030-133-2
US-08-033-401-4
US-08-423-401-4
US-08-423-56-60
US-08-422-964-37
US-08-422-964-37
US-08-65-66-11
US-09-655-832-83
US-09-655-831-10
US-09-655-832-11
US-09-655-837-11
US-09-655-837-11
US-09-655-837-11
US-09-655-837-11
US-09-655-837-11
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US-09-655-837-11
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US-10-121-857-6
US-07-657-769B-58
                                                                                                                                                                                                                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                       513545 segs, 74649064 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                    - protein search, using sw model
                                                                                                                                                                                                                                                                                                                Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query
Match Length DB
                                                                                                                                                                                                                                                                                                                                                                                                                                       Minimum DB seq length: 0 Maximum DB seq length: 4
                                                                                                                                                                                                                                                    1 fglm 4
                                                                                                                                                                                                                                                                                             BLOSUM62
                                                                                                                                                                                                                                                                                             Scoring table:
                                                                                                                                                                                                                              Perfect score:
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                                                                                  OM protein
                                                                                                                                                                                                                                                    Sequence:
                                                                                                                                                                                                                                                                                                                                                     Searched:
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                                                                                                                             Run on:
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No.
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Sequence 63, Application US/08441591

Sequence 63, Application US/08441591

Sequence 63, Application US/08441591

GENERAL INFORMATION:

APPLICANT: NIEDWIANDT, D., GOLD, L. AND WECKER, M. TITLE OF INVENTION: HIGH-AFFINITY

TITLE OF INVENTION: OLIGONUCLEOTIDE LIGANDS

TITLE OF INVENTION: TO THE TACHYKININ

TITLE OF INVENTION: SUBSTANCE P

NUMBER OF SEQUENCES: 66

CORRESPONDENCE ADDRESS:

ADDRESSEE: Swanson & Bratschun, L.L.C.

STREET: Saloue E. Prentice Avenue, Suite 200

CITY: Englewood

STREET: Golorado

COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 MG storage COMPUTER: IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: Wordberfect 5.1
CURRENT APPLICATION DATA:
RPPLICATION NUMBER: US/08/441,591
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435

PRIOR APPLICATION: 435

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/303,362

FILING DATE: 9-SEPTEMBER-1994

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/714,131

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/931,473

FILING DATE: 17-AUGUST-1992

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/117,991

FILING DATE: 8-SEPTEMBER 1993

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/536,428

FILING DATE: 11-AUGUST-1990

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/536,428

FILING DATE: 11-AUGUST-1990

PRIOR DATE: 11-AUGUST-1990

PRIO
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REGISTRATION NUMBER: 33,215
REFERENCE/DOCKET NUMBER: NEX21/C
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 21-OCTOBER-1992
ATTORNEY/AGENT INFORMATION:
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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER REABBLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 MG storage COMPUTER: IBM comparible OPERATING SYSTEM: MS_DOS SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/303,362A
FILING DATE: 9-SEPTEMBER-1994
CLASSIFICATION DATA:
APPLICATION DATA: 735
PRIOR APPLICATION DATA: 77/14,131
FILING DATE: 10-JUNE-1991
PRIOR APPLICATION DATA: 77/931,473
FILING DATE: 17-AUGUST-1992
PRIOR APPLICATION DATA: 8-SEPTEMBER 1993
PRIOR APPLICATION DATA: 1992
FILING DATE: 11-JUNE-1990
PRIOR APPLICATION DATA: 8-SEPTEMBER 1993
PRIOR APPLICATION DATA: 1993
FILING DATE: 11-JUNE-1990
PRIOR APPLICATION DATA: 11-JUNE-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 2
US-08-303-362A-63
i Sequence 62, Application US/08303362A
i Patent No. 5648014
i GENERAL INFORMATION:
i APPLICANT: NIEUWIANDT, D., GOLD, L. AND WECKER, M.
TITLE OF INVENTION: OLIGONUCLEOTIDE LIGANDS
i TITLE OF INVENTION: TO THE TACHYKININ
i TITLE OF INVENTION: SUBSTANCE P
i NUMBER OF SEQUENCES: 66
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Swanson & Bratschun, L.L.C.
STREET: 8400 E. Prentice Avenue, Suite 200
CITY: Englewood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: BATTY J. SWAISON
REGISTRATION NUMBER: 33,215
REFERENCE/DOCKET NUMBER: NEX21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 793-3333
TELEPHONE: (303) 793-3333
TELEFAX: (303) 793-3433
INFORMATION FOR SEQ ID NO: 63:
SEQUENCE CHARACTERISTICS:
LENGTH: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (303) 793-3731 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                         ; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-441-591-63
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TYPE: amino acid
STRANDEDNESS: single
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STREET: 840
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Sequence 2, Application US/09265690C
Fatent No. 6372440
GENERAL INFORMATION:
TITLE OF INVENTION:
TITLE OF INVENTION:
FILE REFERENCE: 1427001
CURRENT FILING DATE: 1999-03-10
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin version 3.0
LENGTH: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Wells, Ibert
APPLICANT: Wells, Ibert
TITLE OF INVENTION: Antagonists of the magnesium binding defect as therapy agents and
TITLE OF INVENTION: methods for treatment of abnormal physiological states
FILE REFERENCE: N1427-002
CURRENT APPLICATION NUMBER: US/09/635,266
CURRENT FILING DATE: 2000-08-09
NUMBER OF SEQ ID NOS: 4
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                                                                100.0%; Score 21; DB 1; Length 4; 100.0%; Pred. No. 4.1e+05; 1ve 0; Mismatches 0; Indels
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100.0%; Score 21; DB 4; L
Best Local Similarity 100.0%; Pred. No. 4.1e+05;
Matches 4; Conservative 0; Mismatches 0;
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Patent No. 6455734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin version 3.0 SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; NAME/KEY: MOD_RES
; LOCATION: (4)...(4)
; OTHER INFORMATION: AMIDATION
US-09-265-6900-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; NAME/KEY: MOD_RES
; LOCATION: (4)...(4)
; OTHER INFORMATION: AMIDATION
US-09-635-266-3
                                                       Query Match
Best Local Similarity 100.
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
Matches 4; Conserv
        1 TOPOLICAL:
US-08-303-362A-63
                                                                                                                                                                                                                                          RESULT 3
US-09-265-690C-2
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Gaps
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Patent No. 5945033

GENERAL INFORMATION:
APPLICANT: YEN, Richard C.K.
TITLE OF INVENTION: NON-CROSSLINKED PROTEIN PARTICLES FOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 4;
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 21; DB 5; L
Pred. No. 4.1e+05;
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                             PRIOR APPLICATION NUMBER: 08/303,362
PRIOR APPLICATION NUMBER: 09-SEPTEMBER-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/361,795
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/117,991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/931,473
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/931,473
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/964,624
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/36,428
FILING DATE: 11-JUNE-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/14,131
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/14,131
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/14,131
PRIOR APPLICATION NUMBER: 07/14,131
PRIOR APPLICATION NUMBER: 07/134,131
PRIOR APPLICATION NUMBER: 07/56,428
FILING DATE: 11-JUNE-1990
ATTORNEY/AGENT INFORMATION:
NAME: BALLY J. SWAIRON:
REFERENCE/DOCKET NUMBER: NEX17/PC
TELECOMMUNICATION:
TELECOMMUNICATION:
                        08/248,632
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TELEFAX: (303) 793-3433
INFORMATION FOR SEQ ID NO: 80:
SEQUENCE CHARACTERISTICS:
                           APPLICATION NUMBER: 08/2/
FILING DATE: 24-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.
.....hes 4, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: single
                                                                        CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: linear
PCT-US95-05600-80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 7
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                                                                                                                                           GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Wells, IDert
TITLE OF INVENTION: Antagonists of the magnesium binding defect as therapy agents and
TITLE OF INVENTION: methods for treatment of abnormal physiological states
TITLE OF INVENTION methods for treatment of abnormal physiological states
TITLE OF INVENTION: Methods for treatment of abnormal physiological states
TITLE OF INVENTION MABER: US/10/230,133
CURRENT FILING DATE: 2002-08-29
FRIOR APPLICATION NUMBER: 09/635,266
PRIOR APPLICATION NUMBER: 09/635,266
PRIOR APPLICATION OF 99
NUMBER OF SEO ID NOS: 4
SOFTWARE: Patentin version 3.0
SEO ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: GOLD, LARRY
APPLICANT: NEUWLANDT, DAN
APPLICANT: WECKER, MATTHEW
APPLICANT: SCHNEIDER, DANIEL J.
APPLICANT: SCHNEIDER, DANIEL J.
APPLICANT: FILGON, JULI
APPLICANT: SULLENERR, BRUCE A.
APPLICANT: DOUDNA, JENNIFER, A.
TITLE OF INVENTION: HIGH-AFFINITY LIGANDS OF
TITLE OF INVENTION: INSULIN RECEPTOR ANTIBODIES, TACHYKININ SUBSTANCE
TITLE OF INVENTION: P, HIV INTEGRASE AND HIV-1 REVERSE TRANSCRIPTASE
NUMBER OF SEQUENCES. 239
CORRESPONDERCE ADDRESS:
ADDRESSED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Swanson & Bratschun, L.L.C.
STREET: 8400 E. Prentice Avenue, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 MG
MEDIUM TYPE: storage
COMPUTER: IBM compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: WORDERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/05600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCT-US95-05600-80
; Sequence 80, Application PC/TUS9505600
; GENERAL INFORMATION:
; APPLICANT: GOLD, LARRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/238,863
FILING DATE: 06-MAY-1994
                                                                                                                Sequence 3, Application US/10230133 Patent No. 6664420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: MOD RES
LOCATION: (4)...(4)
CTHER INFORMATION: AMIDATION
US-10-230-133-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Englewood STATE: Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ouery Match
Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                USA
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                                                                                              US-10-230-133-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGIH: 4
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1 FGL 3
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LENGTH: 3
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US-08-722-126A-20

i Sequence 20, Application US/08722126A

i Patent No. 6034227

i GENERAL INFORMATION:
APPLICANT: BECHT, Israel
APPLICANT: GUTHMANN, Marchel D.
APPLICANT: TAL, Michael
ITILE OF INVENTION: FUNCTION-ASSOCIATED ANTIGEN (MAFA)
CORRESPONDENCES: 20
CORRESPONDENCES: 20
CORRESPONDENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
STREET: 419 Seventh Street N.W., Ste. 300

CITY: Washington
STATE: D.C.
COUNTRY: UNITED STATES OF AMERICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       76.2%; Score 16; DB 2; Length 4;
100.0%; Pred. No. 4.18+05;
Live 0; Mismatches 0; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURENT APPLICATION DATA:
APPLICATION WUMBER: US/08/722,126A
FILING DATE: 08-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                   016197-000840US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; LOCATION; 4
; OTHER INFORMATION: /product= "Met-Amide"
US-08-747-137-124
                             FILING DATE: 12-NOV-1996
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/212,546
FILING DATE: 14-MAR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/069,831
FILING DATE: 01-JUN-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/659,560
FILING DATE: 13-OCT-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/641,720
FILING DATE: 15-JAN-1991
ATTORNEY/ACENT INFORMATION:
NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REPERENCE/DOCKET NUMBER: 016197-0008
TELECOMMUNICATION INFORMATION:
SEQUENCE CHARACTERISTICS:
FINGTH: 4 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/04258
FILING DATE: 06-APR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: Modified-site
CURRENT APPLICATION DATA: APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.
Matches 3; Conservative
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Sequence 2, Application US/09635266
Patent No. 6455734
Fatent No. 6455734
Fatent No. 6455734
FAPPLICANT: Wells, Insert
TILE OF INVENTION: methods for treatment of abnormal physiological states
TITLE OF INVENTION: methods for treatment of abnormal physiological states
CURRENT APPLICATION NUMBER: US/09/635,266
CURRENT FILING DATE: 2000-08-09
NUMBER OF SEQ ID NOS: 4
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; Sequence 2, Application US/10230133
; Sequence 2, Application US/10230133
; Patent No. 664420
; Patent No. 664420
; APPLICANT: Wells, Ibert
; TITLE OF INVENTION: methods for treatment of abnormal physiological states
; TITLE OF INVENTION: methods for treatment of abnormal physiological states
; FILE REFERENCE: 2892-106
; CURRENT APPLICATION NUMBER: US/10/230,133
; CURRENT FILING DATE: 2002-08-29
; PRIOR FILING DATE: 2000-08-09
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ouery Match 71.4%; Score 15; DB 4; Length 3; Best Local Similarity 100.0%; Pred. No. 4.1e+05; Matches 3; Conservative 0; Mismatches 0; Indels
                                                                                                                                                         PECHT=1A
PRIOR APPLICATION DATA:

APPLICATION NUMBER: IL 109257
FILING DATE: 08 -APR.1994
ATTORNEY/AGENT INFORMATION:
NAME: BROWDY, ROGER L.
REGISTRATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: PECHT=1
TELECOWINICATION INFORMATION:
TELEPHONE: (202) 628-5197
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
TENDERAL 4 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.
Matches 3; Conservative
                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
US-08-722-126A-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
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KESUL.
US-08-433-401-4
; Sequence 4, Application US/08433401
· Parent No. 5872097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Amernick, Burton A. REGISTRATION NUMBER: 24,852 REFERENCE/DOCKET NUMBER: 15 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (202) 331-7111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: US
ZIP: 20036-0088
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                              TELEPHONE: (202) 887-040
TELEFAX: (202) 835-0605
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (202) 223-2596
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
                                                                       INFORMATION FOR SEQ ID NO: 6 SEQUENCE CHARACTERISTICS: LENGTH: 4 amino acids TYPE: amino acid STRANDENESS: single: TOPOLGY: linear MOLECULE TYPE: peptide US-08-070-301-8
                                                                                                                                                                                                                                                                                                                                             Query Match 71.4
Best Local Similarity 100.
Matches 3; Conservative
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                                                            TELEFAX:
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APPLICANT: GRANOTO, HILDEN
APPLICANT: KISHIMOTO, JIRO
APPLICANT: FUKCI, Ohji
TITLE OF INVENTION: ENZYME PARTICIPATING IN C-TERMINAL
TITLE OF INVENTION: AMIDATION, AND METHOD OF PREPARING SAME AND USE THEREOF
CORRESPONDENCE ADDRESS:
ADDRESSEE: Wegner, Cantor, Mueller & Player, P.C.
STREET: 1233 20th Street, N.W.
                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                     71.4%; Score 15; DB 4; Length 3; 100.0%; Pred. No. 4.1e+05; rative 0; Mismatches 0; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
PILING DATE: 24-MAY-1991
CLASSIFICATION DATA:
APPLICATION NUMBER: US/08/070,301
FILING DATE: 15-AUG-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US-76331
FILING DATE: 26-MAR-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US-106412
FILING DATE: 24-APR-1990
PRIOR APPLICATION NUMBER: US-2-06412
FILING DATE: 24-APR-1990
PRIOR APPLICATION NUMBER: US-2-06415
FILING DATE: 24-APR-1990
PRIOR APPLICATION NUMBER: US-2-06475
FILING DATE: US-ANG-1990
ATTONENEY/AGENT INPORMATION:
AMPLICATION NUMBER: US-2-AUG-1990
ATTONEY/AGENT INPORMATION:
ANDEL CALLANDER: US-ANG-1990
ATTONEY/AGENT INPORMATION:
AMAMER: US-3-AUG-1990
ATTONEY/AGENT INPORMATION:
AMAMER: US-ANG-1990
ATTONEY/AGENT INPORMATION:
AMAMER: US-ANG-1990
ATTONEY/AGENT INPORMATION:
AMAMER: US-ANG-1990
ATTONEY/AGENT INPORMATION:
AMAMER: US-3-AUG-1990
ATTONEY/AGENT INPORMATION:
AMAMER: US-3-AUG-1990
ATTONEY/AGENT INPORMATION:
AMAMER: US-3-AUG-1990
ATTONEY/AGENT INPORMATION:
AMAMER: US-3-AUG-1990
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REGISTRATION NUMBER: 31,409
REFERENCE/DOCKET NUMBER: P-450-22830
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 8, Application US/08070301
Patent No. 5871995
GENERAL INFORMATION:
APPLICANT: KAMINUMA, TOSHINO
APPLICANT: FUSE, YUKA
APPLICANT: FUSE, YUKA
APPLICANT: TAJIMA, Masahiro
APPLICANT: YAMAGI, Mitsuo
APPLICANT: KISHIMOTO, Jiro
APPLICANT: KISHIMOTO, Jiro
APPLICANT: FUSE, Ohii
NUMBER OF SEQ ID NOS: 4'
SOFWARE: Patentin version 3.0
SEQ ID NO 2
LENGTH: 3
                                                                                                                                                                                 NAME/KEY: MOD_RES
LOCATION: (3)...(3)
CTHER INFORMATION: AMIDATION
US-10-230-133-2
                                                                                                                                                                                                                                                                                                                  Query Match 71.4
Best Local Similarity 100.
Matches 3; Conservative
                                                                                                     TYPE: PRT
ORGANISM: Homo sapiens
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FRY: U.S.A.
20036-8218
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US-08-070-301-8
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APPLICANT: Fryklund, Linda
APPLICANT: Larsson, Bo C.
APPLICANT: Nyberg, Fred J.
APPLICANT: Nyberg, Fred J.
APPLICANT: Ludin, Romny
ITILE OF INVENTION: New Oligopeptides with Affinity to
TITLE OF INVENTION: Opioid Receptors
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPOURE: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/433,401
FILING DATE: 18-MAY-1995
CLASSIFICATION NUMBER: PCT\SE93\00986
FILING DATE: 18-NOV-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: BCT\SE93\00986
FILING DATE: 18-NOV-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: SE 9203496-6
FILING DATE: 20-NOV-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                   71.4%; Score 15; DB 2; Le 100.0%; Pred. No. 4.1e+05; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Pollock, Vande Sande & Priddy STREET: 1990 M Street, N.W., Suite 800 CITY: Washington STATE: D.C.
TELECOMMUNICATION INFORMATION:
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                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: GICCUEL, Brigitte
APPLICANT: GICCUEL, Brigitte
APPLICANT: LIM, Eng Mong
APPLICANT: DATA PORTNO!
DAPLICANT: PERTNO!
APPLICANT: TIMM, Juliano
TITLE OF INVENTION: EXPRESSION VECTORS
NUMBER OF SEQUENCES: 63
CORRESPONDENCE ADDRESS:
ADDRESSE: C/O FINNEGAN, HENDERSON, FARRABOW, GARRETT &
ADDRESSE: DUNNER, L.L.P.
STREET: 1300 I Street, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
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        71.4%; Score 15; DB 2; Length 4; 50.0%; Pred. No. 4.1e+05; tive 2; Mismatches 0; Indels
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CURRENT APPLICATION DATA:

APPLICATION NUMBER:
FILING DATE:
CLASSIFICATION:
APPLICATION NUMBER:
FILING DATE:
FILING DATE:
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FILING APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
APPLICATION NUMBER:
FILING DATE:
APPLICATION NUMBER:
FILING DATE:
APPLICATION NUMBER:
FILING DATE:
APPLICATION NUMBER:
ATORNEY/AGENT INFORMATION:
NAME:
MCGISTERATION NUMBER:
NEGISTERATION NUMBER:
NEGISTERATION NUMBER:
NEGISTERATION
NUMBER:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25, Application US/08793701
Patent No. 6248581
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (202) 408-4132
TELEFAX: (202) 408-4400
INFOTENATION FOR SEQ ID NO. 25.
SEQUENCE CHARACTERISTICS:
Query Match
Best Local Similarity 50.0
Matches 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: protein US-08-793-701-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amino acid
OGY: linear
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1 YGLL 4
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US-08-793-701-25
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RESULT 14 US-09-579-264-25 ; Sequence 25, Application US/09579264

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APPLICANT: GICQUEL, Brigitte
APPLICANT: LIM, Eng Mong
APPLICANT: LIM, Eng Mong
APPLICANT: DIM, Eng Mong
APPLICANT: TIMM, Juliano
TITLE OF INVENTION: EXPRESSION VECTORS
NUMBER OF SEQUENCES: 63
CORRESPONDENCE ADDRESS:
ADDRESSEE: C/O FINNEGAN, HENDERSON, FARRABOW, GARRETT &
STREET: 1300 I Street, N.W.
CITY: Washington
STATE: D.C.
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 37, Application US/08429964

Patent No. 2962243

GENERAL INFORMATION

APPLICANT: BROWN, MICHAEL S.

APPLICANT: GOLDGTEIN, JOSEPH L.

APPLICANT: JAMES, GUY L.

TITLE OF INVENTION: WETHODS FOR THE IDENTIFICATION OF FARNESYL

TITLE OF INVENTION: TRANSFERASE INHIBITORS

CORRESPONDENCES: 85

CORRESPONDENCE ABDRESS:

ADDRESSEE: ADDRESSE:

ADDRESSEE: P.O. BOX 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14; DB 4; Length 4;
Pred. No. 4.1e+05;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PATENTIN Release #1.0, Version #1.30 APPLICATION NUMBER: US/09/579,264 FLING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: MCDONell, Lealie A.
REGISTRATION NUMBER: 34,872
REFERENCE/DOCKET NUMBER: 02356.0075
TELECOMMUNICATION:
TELEPHONE: (202) 408-4132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/793,701
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: FR 94/10585
FILING DATE: 02-SEP-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (202) 408-4400
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       66.78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 66.7
Matches 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 4 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: HOUSTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-579-264-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 15
US-08-429-964-37
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Search completed: March 23, 2005, 15:14:28 Job time : 42 secs

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